

1 **Title:** Antimicrobial activity and regulation of CXCL9 and CXCL10 in oral
2 keratinocytes.

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14 **Running head:** CXC ELR- chemokines in oral mucosa

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28 keratinocytes.

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30

31 **Abstract**

32 It is well established that the CXCL9 and CXCL10, two chemokines that belong to the
33 CXC family, are dysregulated in oral inflammatory conditions. It has not been
34 established if these chemokines target the micro-organisms that form the oral biofilm.
35 The aim of this study was thus to investigate the antimicrobial activity of CXCL9 and
36 CXCL10 on oral microflora and their expression profile in oral keratinocytes following
37 inflammatory and infectious stimuli. *Streptococcus sanguinis* was used as a model and
38 *Escherichia coli* as a positive control. The antimicrobial effect of CXCL9/CXCL10 was
39 tested using radial diffusion assay. mRNA transcripts were isolated from LPS-treated
40 and untreated (control) oral keratinocyte cell lines at 2, 4, 6 and 8 h. The CXCL9/10
41 expression profile in the presence or absence of IFN- γ was assessed using semi-
42 quantitative PCR. Both chemokines demonstrated antimicrobial activity. CXCL9 was
43 the most effective against both *Strep. sanguinis/Ecoli*. CXCL10 mRNA was expressed
44 in control cells and its production was enhanced by LPS stimulation at all time-points.
45 Conversely, CXCL9 mRNA was not expressed in control or LPS-stimulated cells.
46 Finally, IFN- γ stimulation enhanced basal expression of both CXCL9 and CXCL10 in
47 oral keratinocytes. Oral epithelial derived chemokines demonstrate antimicrobial
48 properties, particularly CXCL9. Bacterial and inflammatory stimulated up-regulation
49 of CXC9/10, with the potential to prevent adherence and invasion of oral epithelial

50 cells, could represent a key element in oral bacterial colonization homeostasis and host
51 defense mechanisms.

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53 **Keywords:** Chemokines; CXCL10; CCL28; XCL9; Streptococci sanguis.

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59

60 **Introduction**

61 Bacteria colonize all surfaces of humans, but are particularly dense in lower
62 gastrointestinal tract and in the oral cavity where Streptococci represent a large
63 proportion of the resident microflora.

64 It has previously been shown that bacteria can adhere and invade host oral epithelial
65 cells (1) and furthermore, there are bacterial receptors present in saliva that can be
66 absorbed onto oral mucosal surfaces (2). Among them, the toll like receptors (TLR)
67 expressed on host cells are involved in the recognition of conserved bacterial patterns,
68 for example the cell wall component of gram negative bacteria, lipopolysaccharide
69 (LPS) (3).

70 In order to counter bacterially-mediated tissue damage, keratinocytes produce a wide
71 number of molecules encompassing antimicrobial peptides (AMPs), that are
72 differentially regulated together with pro-inflammatory cytokines across epidermal
73 layers following bacterial stimuli (4).

74 The chemokines monokine induced by IFN- γ CXCL9, and IFN- γ -induced protein-10
75 CXCL10 are two chemokines belonging to the CXC family, both binding the same
76 CXCR3 receptor (5). Chemokines play an important role in directing the migration of
77 specific immune cell populations and for some a direct anti-bacterial and/or -fungal
78 activities has been demonstrated (6-8). As oral epithelial cells are known to produce
79 chemokines (9-11) these may be playing a direct role in microbial defence.

80 It is known that the CXC ELR- chemokines can be induced by LPS in some cell types
81 (12), and that these chemokines are expressed during bacterial infections (13). It is also
82 known that LPS is capable of inducing a range of cytokines and chemokines from
83 epithelial cells predominately by signalling through TLR-4 (14).

84 The induction of these chemokines may be triggered in response to an alteration in the
85 microbial flora, which could in turn cause an ensuing immune cell infiltration. We
86 recently found that CXCL9/10 are dysregulated in oral inflammatory disease (personal
87 communication), but nothing is known about their antibacterial activity. Furthermore,
88 while the induction of several cytokines in oral epithelial cells after LPSs stimulation
89 was previously demonstrated (11,14-17), the role of CXCL9/10 in the infection and
90 immunity of the oral cavity has never been investigated.

91 Therefore the aim of this paper was to investigate the potential of CXC ELR-
92 chemokines to mediate microbicidal activity on the gram positive *Strep. sanguinis*, one
93 of the most prevalent residents of the oral microflora, and the chemokines expression
94 in oral keratinocytes after exposure to infectious and inflammatory stimuli.

95 The CCR10 ligands CCL27 (CTACK) and CCL28 (MEC) are two C-C chemokines
96 that bind the CCR10 receptor found to exert a potent antimicrobial activity
97 against *Candida albicans*, Gram-negative bacteria, and Gram-positive bacteria (7). We
98 used these two chemokines as positive controls.

99

100 **Material and Methods**

101

102 *Cell culture techniques*

103 *Normal Human Oral keratinocytes (NHOK) cell culture.*

104 All normal oral mucosa was obtained from healthy patients attending the Oral Surgery
105 Clinic, Eastman Dental Institute for routine third molar extraction. Three different
106 NHOK strains (NHOK1, NHOK2, NHOK3) were isolated from the excised normal
107 tissue. The samples were cut into approximately 1mm³ pieces and culturing at 37°C
108 /5% CO₂ in keratinocyte basal medium-2 containing the recommended growth
109 supplements (Biowittaker, Wokingham, UK). The epithelial cells were then detached
110 using 0.25% trypsin - 1 mM EDTA. The viability of the keratinocytes was confirmed
111 by trypan blue exclusion. All of the cell lines/strains were derived prior to 2001 and
112 therefore were not subject to ethical committee approval in the UK (18). The study was
113 approved by the internal research committee at Eastman Dental Institute, University
114 Collage London.

115

116 *H357 cell culture*

117 The oral squamous cell carcinoma cell line, H357, was established by PRIME *et al* (19).
118 This cell line was grown in the same medium as described for the NHOK.

119

120 *Bacterial cell culture and antimicrobial assessment*

121 All bacterial stocks were maintained frozen at -70°C in trypticase soy broth (TSB)
122 (Becton Dickinson) supplemented with 0.6% yeast extract (YE) (Oxoid,
123 Basingstoke,UK) and 10% glycerol (BDH). Cultures were checked weekly both

124 visually and by Gram-staining for contamination with other bacteria. Stocks of *E. coli*
125 NCTC JM22 and *Strep. sanguinis* NCTC 10904 (provided by Dr.Rod McNab at the
126 Eastman Dental Institute, University College London, UK) bacterium were plated upon
127 an agar plates containing 3% TSB. They were grown for 48 h at 37°C / 5% CO₂ and
128 maintained by twice weekly subculture on TSB agar plates.

129 Each *E. coli* colony was taken and resuspended into 50 ml of 3% TSB. For the oral
130 streptococci species 3 streaks of a cultured plate were resuspended into 10 ml of 3%
131 TSB. The cultures were shaken at 250 rpm at 37°C for 15-18 h. 50 ml of the *E. coli*
132 culture or 2 ml of the oral streptococci species culture was transferred into 50 ml or 10
133 ml of 3% TSB respectively. This was shaken at 250 rpm for 3.5 h at 37°C. After this
134 time the culture was adjusted to an OD of 1 at 620 nm.

135 To prepare the underlay, 50 ml of 100 mM sodium phosphate buffer, 5 g agarose, low
136 EEO (Sigma, Poole, UK) and 5 ml 3% trypticase soy broth (TSB) were added to 1 litre
137 of distilled water. Then the pH was then adjusted to 7.4, the agarose was dissolved by
138 heating the solution in the microwave. Then the solution was dispensed into in 50 ml
139 aliquots and autoclaved. The underlay aliquots were then stored at room temperature
140 until use in the radial diffusion assays at which point they were heated in a microwave
141 until fluid and then stored in a 60°C water bath.

142 8 ml of *E. coli* or 16 ml of the streptococcal species was added to 5 ml of molten
143 underlay and dispensed into a Petri dish, using a leveling tray. This underlay was
144 allowed to set, then 3 mm holes were punched in the gel, using 10 ml pipettes
145 (Starstedt). 5 µl of test solution diluted in 0.01% acetic acid was added to the wells.

146 10 g agarose, low EEO (Sigma, Poole, UK) was added to 6% TSB, aliquoted into 50
147 ml and autoclaved. The overlay aliquots were then stored at room temperature prior the
148 use in the radial diffusion assays.

149 This plate was incubated for 3 h at 37°C, before 5 ml of overlay was added to the plate
150 and incubated at 37°C overnight.

151 *Radial diffusion assays* were then performed, adding 5 µl of either recombinant human
152 CXCL9/CXCL10/CCL27/CCL28 (all Peprotech EC, London, UK) or 0.01% acetic
153 acid to the wells before incubating the plates. 100 µM tetracycline was used as a
154 positive control for the assay.

155 Images of the plates were taken using Alphaimager software and the zones around the
156 cultures measured from 3 different points from the end of the well.

157

158 *IFN-γ cell treatment assay*

159 In a modification of the method utilised by ALTENBURG *et al* (20), the H357 cells were
160 seeded at 8×10^4 cells / well in a Falcon 6 well plate (Becton Dickinson, Oxford, UK)
161 with 3 ml of KBM-2 medium containing no hydrocortisone. The cells were incubated
162 for at least 3-5 days until cell culture was 60-80% confluent. Medium containing 1000
163 U ml⁻¹ IFN-γ was added to 3 wells and control cell culture medium only was added to
164 the remaining 3 wells. The cells were incubated for 48 h. The supernatant was extracted,
165 centrifuged and stored at -70°C. The adherent cells were washed with PBS (Gibco Life
166 Technologies, Paisley, UK) before 0.5 ml of Trizagent (Sigma, Poole, UK) were
167 added. The suspension was then removed and stored at -70°C. The RNA was isolated
168 as described below.

169

170 *CXCL9 and CXCL10 mRNA transcripts in oral epithelial cells in response to*
171 *lipopolysaccharide (LPS) and IFN- γ : mRNA isolation and semi-quantitative RT-*
172 *PCR*

173 mRNA transcripts for 18S, CXCL9 or CXCL10 were isolated from H357 cells without
174 (CON) or treated with LPS (LPS) for 2, 4, 6 or 8 h. The RNA was extracted using TRI
175 Reagent® (Sigma, Poole, UK) and 2 ml Pellet Paint Co-precipitant (Novagen,
176 Nottingham, UK) to visualise the RNA pellet.

177 Single strand cDNA synthesis was performed. 2 ml of RNA was added to 4 ml
178 deoxynucleotides (dNTPs) (2.5 mM) (Sigma), 2 ml of random hexamers (50 mM)
179 (Ambion, Texas, USA) and 9.5 ml dH₂O. Then 1 ml of RNAaseIN (Ambion, Austin,
180 Texas, USA), 2 ml 10x MuLVRT buffer and 0.5 ml M-MuLVRT (200 U ml⁻¹)
181 (Boehringer-Mannheim, Germany) was added and incubated at 42°C for 1 h.

182 RT-PCR for 18S, CXCL9 and CXCL10: Magnesium concentration was optimised for
183 each primer as follows; 1 ml of cDNA was added to 4 ml dNTP (2.5 mM), 5 ml 10x
184 buffer, 0.225 ml AmpliTaq (5.0 U ml⁻¹) (Perkin Elmer), 4 ml of each specific primer
185 (5 mM), 1.5 mM, 3.0 mM or 4.5 mM MgCl in each reaction and dH₂O added to give a
186 final volume of 50 ml. The products were separated on a 2% agarose (GibcoBRL Life
187 Technologies, Paisley, UK) gel and visualised by staining with ethidium bromide
188 (Sigma, Poole, UK), specific bands were visualised by ultra-violet trans-illumination
189 in a MultiImage Light Cabinet (AlphaInnotech Corp., Cannock, UK) and digital images
190 acquired and stored using AlphaImager Software (AlphaInnotech Corp., Cannock,
191 UK).

192 CXCL9 (5'-ccaacacccccagaagtgc-3', 5'-gccagcacctgctctgagac-3'), CXCL10 (5'-
193 gccaatgttccacgtgttg-3', 5'-aaagaattggggccccttg-3'), and 18S ribosomal RNA (5'-
194 ttccggaactgaggccatga-3', 5'- gcatgccagagtctcgttcg -3') primers were generated for use

195 in this study (Genosys-Sigma, Poole, UK). The thermocycler (Techne Genius;
196 Cambridge, UK) parameters utilised were at 94°C for 45 s, 57°C for 45 s, 72°C for 45
197 s.

198 For each primer the linear range was determined by repeating the above reaction with
199 optimised magnesium concentration for each primer and stopping the reaction at 15,
200 17, 19, 21, 23, 25, 27, 29, 31, 33 and 35 cycles. The mid-point of each linear range was
201 determined by using intensity analysis of the bands with AlphaImager software, and
202 this cycle length was utilised for each primer in subsequent reactions. 18S primer and
203 18S Competitor primers (Ambion, Texas, USA) were combined to ratios 1:9, 2:8 and
204 3:7 respectively. For each of the primers CXCL9/10, 4 ml of the 18S primer: competitor
205 mix was added to the RT-PCR reaction. The band intensity of the 18S and specific
206 primer was quantified for each primer in each sample with Phoretix 1D software
207 (Phoretix, Newcastle, UK).

208 Unless otherwise specified all the experiments were performed at least in triplicate.

209

210

211 **Results**

212 *CXCL9 and CXCL10 mRNA transcripts in an oral epithelial cell line in response to*

213 *IFN- γ*

214 The production of CXCL9/10 chemokines in oral mucosal keratinocytes was first
215 assessed over time in preliminary experiments using the keratinocyte cell line H357
216 (Supplementary Figure 1). CXCL10 mRNA levels in H357 cells were detected as early
217 as 3 h after IFN- γ treatment and appeared to peak at 24 h. In contrast, the control cells
218 showed virtually undetectable mRNA levels over the same time period. CXCL9 mRNA
219 transcripts demonstrated a similar pattern to CXCL10, with a biphasic pattern showing

220 a rapid induction of mRNA in the stimulated cells followed by a second peak at 24/48
221 h. Thus, the expression of CXCL9/10 in H357 cells can be significantly enhanced by
222 IFN- γ in a time-dependent manner, with a peak after 48 h (Figure 1). Similar results
223 were obtained in primary normal oral keratinocytes (NHOK) (Figure 2). These data
224 were confirmed at the protein level by ELISA (data not shown). These results show that
225 the expression of CXCL9/10 in oral epithelial cells are significantly enhanced by IFN-
226 γ in a time-dependent manner.

227

228 *CXCL9 and CXCL10 mRNA transcripts in an oral epithelial cell line in response to*
229 *LPS*

230 CXCL10 mRNA was expressed in control cells (without LPS exposure), but was
231 enhanced in response to LPS stimulation after 2, 4 and 6 h incubation. This expression
232 subsequently decreased after 8 h of stimulation (Figure 3). In contrast, CXCL9 mRNA
233 was not expressed in either control or LPS-stimulated cells at any of the time-points
234 tested.

235

236

237 *Antimicrobial effect of CXCL9, CXCL10, CCL27 or CCL28 upon Strep. sanguinis and*
238 *E. coli*

239 CCL27 and CCL28 were used as positive controls. The clear zones of bacterial growth
240 depletion that were produced after antimicrobial activity of the chemokines are
241 indicated in Figure 4 and Figure 5. All of the chemokines tested demonstrated a level
242 of antimicrobial activity at the tested concentration (Table 1). CXCL9 was the most
243 effective of the tested chemokines against both *Strep. sanguinis* and *E. coli*. CCL27 and
244 CXCL10 had a less effective antimicrobial action against *Strep. sanguinis* compared to

245 *E.coli*. CCL28 had the greatest antimicrobial action against *E. coli*. These results clearly
246 demonstrate that the epithelial-derived chemokines CXCL9/10 exert antimicrobial
247 activity.

248

249 **Discussion**

250 Chemokines are known to have antimicrobial effects but little is known of the action of
251 chemokines derived from the oral mucosal epithelium. The present study is the first to
252 examine the expression of CXC ELR- chemokines by oral epithelial cells when
253 stimulated by bacteria-derived products such as LPS. The present study has established
254 that a cell line of oral origin is capable of expressing mRNA of CXCL9/CXCL10 when
255 stimulated with LPS. The production of these chemokines was also enhanced by
256 inflammatory stimuli such as IFN- γ stimulation.

257 CXCL10 mRNA expression was increased in H357 cells after LPS stimulation,
258 whereas CXCL9 mRNA expression was not induced over the same time-period.
259 Previously we have shown that CXCL10 can act as a potent chemoattractant of
260 lymphocytes, hence this local production of CXCL10 by oral epithelial cells in response
261 to LPS could have important effects upon oral inflammation, perhaps crucially in the
262 initial stages of inflammation.

263 Previous studies have also shown that LPS treatment alone can induce or enhance
264 CXCL10 mRNA expression in several different cell types (12,21-23). However, in
265 contrast to the present findings, LPS stimulation did not induce CXCL10 from cultured
266 skin keratinocytes (24), suggesting perhaps that oral keratinocytes are more responsive
267 to LPS stimulation than cutaneous keratinocytes. This difference in expression may
268 reflect the high bacterial load in the oral cavity.

269 The rapid expression of mRNA of CXCL10 observed in the present study accords with
270 that of murine macrophages (17,25) although LPS stimulation may be more transient
271 than IFN- γ in CXCL10 production (10). This short-term effect may be essential to avoid
272 over-stimulation of CXCL10 in response to resident bacteria in proximity to the
273 epithelium. IL-10 is known to be able to down-regulate production of LPS-induced
274 CXCL10 in macrophages (22). This cytokine is present within oral lichen planus (OLP)
275 lesions and is increased in serum and saliva from patients with oral lichen planus (26),
276 and thus may act to down-regulate expression of CXCL10 in oral inflammation.

277 There are few reports of LPS-induced production of CXCL9. The present study
278 revealed that CXCL9 mRNA was not expressed by H357 cells when stimulated with
279 LPS. In contrast to the present study, CXCL9 mRNA was found to be expressed in
280 LPS-stimulated murine dendritic cells (12), however, in another study, this chemokine
281 was not induced in the same murine cell line by LPS, despite induction of CXCL9 by
282 IFN- γ (25).

283 It is possible that CXCL9 displays a delayed response in comparison to CXCL10, as
284 CXCL9 mRNA induction in lung tissue of intravenously LPS-treated mice displays a
285 later induction to CXCL10, and is never expressed to the same levels as CXCL10 (25).

286 Many studies report that LPS and IFN- γ act synergistically to induce the production of
287 high levels of CXCL10 mRNA, for example, in breast carcinoma cells (21,22). It is
288 then possible that CXCL10 levels could be enhanced in oral inflammation where there
289 is a presence of both LPS and IFN- γ , perhaps through the enhancement of specific
290 TLRs (27).

291 Gram-positive bacteria, such as *Strep. sanguinis* contain components other than LPS
292 that are known to stimulate chemokine release from various cell types and it would be
293 interestingly to determine whether these are also capable of stimulating the production

294 of the CXC ELR- chemokines in oral epithelium. Many gram-positive bacterial
295 components act on a different toll like receptor, TLR-2, which is functionally expressed
296 upon keratinocytes (28). However, TLR-2 agonists do not induce CXCL 10 production
297 in macrophages (29) or dendritic cells (30) in vitro. This suggests that TLR-2 agonistic
298 bacterial products would also not induce CXCL10 in epithelial cells. Therefore, only
299 products bound by TLR-4 would be influential in upregulating CXCL10 production in
300 epithelial cells. In addition, as LPS-mediated CXCL10 production is TLR-4 dependant,
301 it strongly suggests that oral keratinocytes bear functional TLR-4, therefore, LPS-
302 stimulation of epidermal cells is not due to TLR-2 agonist contaminants in LPS
303 preparations, as previously suggested (28).

304 Chemokine modulation in oral cells by bacterial products is thus complex and many
305 different factors, including T cell contact, may play a factor in chemokine induction
306 during an immune response.

307 The present studies suggest that in certain circumstances, bacterial products could
308 stimulate oral epithelial cells to produce an inflammatory response through TLR-4
309 agonists (perhaps after continuous TLR-2 stimulation) and this may induce epithelially-
310 derived CXCL10-mediated inflammation. This inflammation would presumably be
311 characterised by activated memory T cell infiltration localised under the basal
312 epithelium reminiscent of the pathology of OLP.

313 CXCL9 was shown to be a potent antimicrobial agent against both *Strep. sanguinis* and
314 *E. coli*, as did CXCL10 but to a lesser degree, confirming the study by COLE et al (6).

315 Although only one oral bacterial species was tested in the present study, the data hints
316 that antimicrobial properties of these chemokines may assist in countering bacterial
317 growth in the oral cavity.

318 The choice of using *S. sanguinis* as a model has been driven by the delicate role that
319 this bacterium plays in the maintenance of the oral flora balance. It is commonly found
320 in healthy tissues as pioneer colonizer as well as it is implicated in modulating the
321 virulence of bacterial biofilms (31). Furthermore, significant inhibitory effects of the
322 intracellular proteins produced by *S. sanguinis* on the growth and the morphology of
323 many other components of the oral flora, such as *P. intermedia*, *P. gingivalis*, *C.*
324 *albicans*, *C. tropicalis* and their biofilms have been demonstrated (32,33). This allow
325 us to conjecture and expect a series of chain effects if the CXCL9/10 antimicrobial
326 activity is transferred in an *in vivo* model. As previously showed for the two
327 chemokines CCL27 and CCL28, for which a wide spectrum of antimicrobial activity is
328 well established, the action of CXCL9/10 expected against other micro-organisms of
329 the oral cavity should follow a similar pattern (34).

330 The low production of CXCL9 by oral epithelial cells following IFN- γ or LPS
331 stimulation could potentially be a means of avoiding an over-active anti-microbial
332 response due to the potent antimicrobial activity of CXCL9.

333 In our study both CCL27/CCL28 showed to exert antimicrobial properties against *E.*
334 *coli* and *Strep. sanguinis*. CCL28 has been shown to have a microbicidal activity
335 against a wide range of bacteria (both Gram-negative and Gram-positive) and yeasts
336 (7, 35), and our findings confirmed that this chemokine was also effective against the
337 oral commensal *Strep. sanguinis*. This is only the second report of the antimicrobial
338 effect of CCL28 upon the enterobacterial *E. coli* after the study reported by BERRI et al
339 in 2014 (35).

340 The production of CXCL10 by oral epithelial cells, in inflammatory conditions such as
341 OLP, may be induced by resident bacteria that contain TLR-4 agonists. If this
342 inflammation resulted from constant TLR-2 stimulation, this may cause a CXCL10

343 based inflammation and a resultant influx of memory Th1 CD4+ cells. Furthermore,
344 CXCL10 is antimicrobial at high concentrations. It may be upregulated by the presence
345 of bacteria, thus playing a role in defense of oral epithelial cells. The antimicrobial
346 activity of the chemokines tested is likely to be an important mechanism in the
347 homeostasis of oral bacterial colonisation. Any IFN- γ in the oral epithelial area (which
348 could presumably be produced by the infiltrating Th1 cells) may synergise with TLR-
349 4 agonists to cause an increased inflammatory state. Further studies are still warranted
350 to confirm these novel findings.

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488 **Figure legends:**

489 **Figure 1:** 18S, CXCL9 and CXCL10 mRNA expression in the H357 cell line either
490 interferon-gamma treated for 48 h (IFN) or control cells that were left untreated (CON)
491 over the same time period. The left column “m.w.” indicates the molecular weight
492 (marker column).

493

494 **Figure 2:** CXCL9 and CXCL10 mRNA expression in 3 different normal human oral
495 keratinocytes (NHOK1, NHOK2, NHOK3) either interferon-gamma treated for 48 h
496 (IFN) or control cells that were left untreated for 48 h (CON). The values were
497 normalized against 18S mRNA expression used as house keeping.

498

499 **Figure 3:** a) CXCL9 and CXCL10 expression by H357 cells with or without LPS
500 stimulation: mRNA transcripts for 18S, CXCL9 or CXCL10 in H357 cells without
501 (CON) or treated with LPS (LPS) for 2, 4, 6 or 8 h. b) Densitometric analysis of
502 CXCL10 mRNA relative to 18S expression with (grey) and without (blue) LPS
503 treatment.

504

505 **Figure 4:** Radial diffusion gels demonstrating the antimicrobial activity of $100 \mu\text{g ml}^{-1}$
506 human recombinant CXCL9, CXCL10, CCL27 and CCL28 upon *Strep. sanguinis* and
507 *E. coli* bacterium. The negative control (-ve) contains 0.01% of acetic acid only. The
508 diameter of the cleared zone around the well containing chemokine represents the
509 antimicrobial properties of the chemokine.

510

511 **Figure 5:** The antimicrobial activity of CXCL9 on *E. coli* JM22 bacterium at 50, 10,
512 and 1 μM . 0 μM contains 0.01% of acetic acid only.

513

514 **Supplementary Figure 1:** 18S, CXCL9 and CXCL10 mRNA expression in the H357
515 cell line. (ifn)= cells that were treated with interferon-gamma for 3, 6, 9, 24, 48 or 72
516 h; (con)= control cells that were left untreated over the same time period.

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