



Table S1. Demographic characteristics and periodontal (PDL) health status of the study participants.

		Control (n=24)	CC_MC (n=35)	CC-Zr (n=35)	MC (n=35)	P-value
Gender n (%)	Male	11 (45.8%)	17 (48.6%)	16 (45,75)	17 (48.6%)	P = 0.11
	Female	13 (54.2%)	18 (51.4%)	19 (54.3%)	18 (51.4%)	P = 0.14
Age (years)	Range	19-47	18-48	19-49	20-50	P>0,5
	Mean±SD	34.8±7.8	32.3±8.5	35.1±6.8	36.6±8.2	
PDL status		Healthy periodontium, no bleeding/periodontal pockets	Clinical examination confirmed the diagnosis of periodontitis in patients			

Table S2. Multiple comparison of the microbiological composition and counts of the gingival crevicular fluid in all groups before and after treatment.

Microorganisms	Before treatment					
	Healthy vs MC	Healthy vs CC-MC	Healthy vs CC-Zr	MC vs CC-MC	MC vs CC-Zr	CC-MC vs CC-Zr
<i>Enterococcus spp.</i>	0.741	0.976	0.381	0.940	0.883	0.624
<i>Peptostreptococcus spp.</i>	0.980	0.927	0.976	0.993	0.844	0.738
<i>Neisseria spp.</i>	0.979	0.041	0.966	0.065	0.999	0.149
<i>Peptococcus spp.</i>	1	0.683	1	0.553	0.998	0.735
<i>Staphylococcus spp.</i>	1	0.932	0.004	0.923	0.002	0.023
<i>Beta-hemolytic streptococcus</i>	0.362	0.198	1	0.960	0.394	0.221
<i>Candida albicans</i>	0.600	0.977	1	0.846	0.650	0.985
<i>Alpha-haemolytic streptococcus</i>	0.996	0.058	0.808	0.060	0.878	0.400
<i>Lactobacillus spp.</i>	0.995	0.998	0.877	0.970	0.721	0.939
<i>Corynebacterium spp.</i>	0.830	0.015	1	0.078	0.803	0.015
<i>Fusobacterium spp.</i>	0.636	0.901	1	0.636	0.970	0.907
<i>Porphyromonas gingivalis</i>	1	1	0.208	1	0.167	0.207
<i>Prevotella intermedia</i>	1	1	0.285	1	0.229	0.280
Microorganisms	At 6 months of treatment					
	Healthy vs MC	Healthy vs CC-MC	Healthy vs CC-Zr	MC vs CC-MC	MC vs CC-Zr	CC-MC vs CC-Zr
<i>Enterococcus spp.</i>	0.065	0.982	0.992	0.153	0.154	1
<i>Peptostreptococcus spp.</i>	0.653	0.871	0.930	0.189	0.283	0.999
<i>Neisseria spp.</i>	0.735	0.494	0.631	0.961	0.992	0.998
<i>Peptococcus spp.</i>	0.949	0.525	1	0.181	0.958	0.532
<i>Staphylococcus spp.</i>	0.266	0.995	0.657	0.152	0.952	0.497
<i>Beta-hemolytic streptococcus</i>	0.025	1	1	0.024	0.31	1
<i>Candida albicans</i>	0.408	1	1	0.397	0.431	1
<i>Alpha-haemolytic streptococcus</i>	0.968	0.427	1	0.637	0.983	0.492
<i>Lactobacillus spp.</i>	0.833	0.973	0.724	0.980	0.990	0.921
<i>Corynebacterium spp.</i>	0.729	0.980	0.923	0.923	0.333	0.743
<i>Fusobacterium spp.</i>	0.936	0.650	0.589	0.248	0.217	0.999
<i>Porphyromonas gingivalis</i>	0.645	1	0.966	0.692	0.921	0.979
<i>Prevotella intermedia</i>	0.657	0.994	0.271	0.812	0.842	0.395
<i>Veillonella spp.</i>	0.759	0.997	0.271	0.864	0.755	0.362
<i>Corynebacterium anaerobium</i>	0.740	0.571	0.999	0.983	0.825	0.667

At 12 months of treatment						
<i>Enterococcus spp.</i>	0.064	0.997	0.668	0.099	0.631	0.774
<i>Peptostreptococcus spp.</i>	0.318	0.207	0.491	0.207	0.972	0.972
<i>Neisseria spp.</i>	0.927	0.968	0.113	0.999	0.265	0.265
<i>Peptococcus spp.</i>	0.224	0.118	0.913	0.964	0.051	0.025
<i>Staphylococcus spp.</i>	0.150	1	0.957	0.149	0.430	0.961
<i>Beta-hemolytic streptococcus</i>	0.001	0.994	1	0.002	0.002	0.998
<i>Candida albicans</i>	0.555	1	1	0.535	0.565	1
<i>Alpha-haemolytic streptococcus</i>	0.411	0.961	0.346	0.731	0.992	0.630
<i>Lactobacillus spp.</i>	0.213	0.211	0.232	0.999	0.999	1
<i>Corynebacterium spp.</i>	0.463	0.431	0.457	0.998	0.998	1
<i>Fusobacterium spp.</i>	0.556	0.871	0.850	0.37	0.138	1
<i>Porphyromonas gingivalis</i>	0.916	0.975	0.593	0.997	0.890	0.830
<i>Prevotella intermedia</i>	0.159	0.186	0.946	1	0.479	0.497
<i>Veillonella spp.</i>	0.180	0.933	0.198	0.501	0.999	0.493
<i>Corynebacterium anaerobium</i>	0.915	0.155	1	0.370	0.923	0.173

Post hoc Scheffe. Significant at $P < 0.05$.