

**Table S1** Barcodes used for different biofilm samples

Description of biofilm samples	Barcode Sequence
Stone 10°C	AAAACACA
Tyre- derived rubber10°C	AAAACGAC
Stone 20°C	AAAACGAG
Tyre -derived rubber 20°C	AAAACACC
Stone 30°C	AAAACGCC
Tyre- derived rubber 30°C	AAAACGCG

**Table S2** Physico-chemical properties of the influent and effluent from aerobic reactors at different temperature conditions for stone (ST) and tire derived rubber (TDR) media

Temperature (°C)	Parameters analyzed	Media material	
		ST	TDR
10	BOD (mg/L)	Influent	320.83 ±13.15
		Effluent	208.91±09.5
	DO (mg/L)	Influent	2.06±0.03
		Effluent	4.83±1.0
	pH	Influent	7.30±0.2
		Effluent	7.24±1.02
20	BOD (mg/L)	Influent	378.09±26.9
		Effluent	278.66±10.12
	DO (mg/L)	Influent	1.95±0.7
		Effluent	5.37±1.1
	pH	Influent	7.86±0.3
		Effluent	7.47±0.6
30	BOD (mg/L)	Influent	320.83±13.15
		Effluent	181.77±10.09
	DO (mg/L)	Influent	2.06±0.03
		Effluent	5.84±1.0
	pH	Influent	7.30±0.2
		Effluent	7.03±0.4

Key = ± Standard Deviation

**Table S3** MOTHUR diversity indices of bacterial communities in twelve aerobic biofilm samples developed on stone (ST) and tire derived rubber (TDR) media

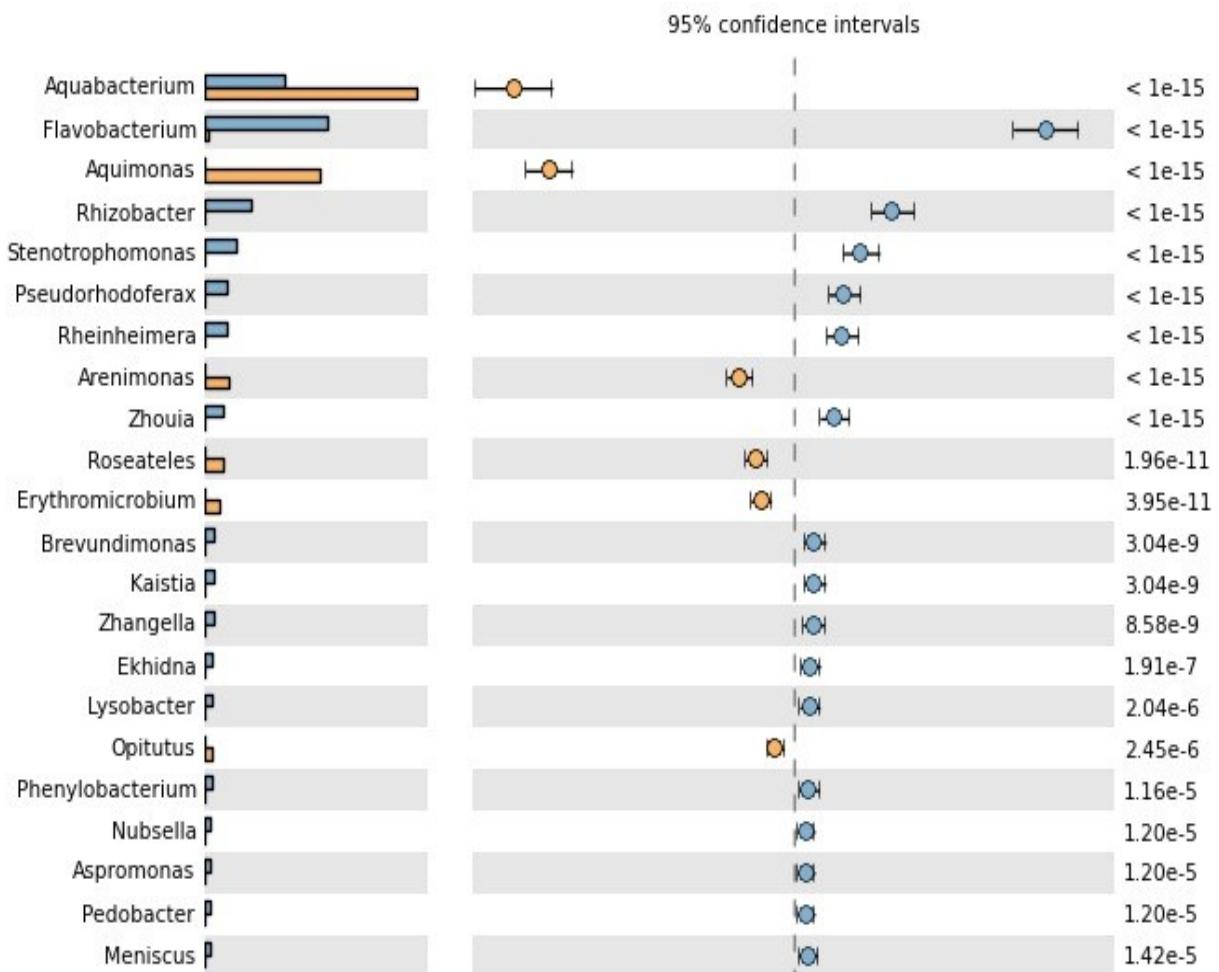
	Nseqs	Sobs	Chao	Chao_Lci	Chao_Hci	Ici	Hci	Ace	Ace_Lci	Ace_Hci	Lci	Hci	Invsimpson	Lci	Hci	Coverage
ST10°C	2004	289	404.9565	362.78	471.2433	29.82329	46.87182	464.4569	410.2067	542.9886	38.36069	55.53034	8.620901	0.598613	0.744912	0.936627
TDR 10°C	1714	270	402.2553	351.0448	485.8247	36.21131	59.09249	485.2294	434.3552	551.851	35.97345	47.10859	42.74893	2.838531	3.494899	0.934656
ST20°C	2272	257	379.9362	331.8071	459.0302	34.03236	55.92792	476.9426	424.696	545.4668	36.9439	48.45396	17.84281	0.986054	1.168737	0.952465
TDR 20°C	1909	347	608.0169	521.3074	737.8604	61.31292	91.81315	823.0073	734.5902	931.5942	62.52035	76.78251	42.70273	2.638934	3.197896	0.907805
ST 30°C	1324	256	383.5532	333.8942	464.8708	35.11423	57.50021	395.6414	349.7696	463.9533	32.4362	48.30385	45.5353	4.121054	5.538898	0.916918
TDR 30°C	2919	266	406.0217	351.999	493.9804	38.19983	62.19614	487.4798	435.1462	556.0053	37.00545	48.45488	9.358689	0.415459	0.475116	0.960946

**Key:** nseqs: number of reads analysed, sobs: species observed at the cut-off value (no of observed OTUs), ace: abundance-based coverage estimator, chao: abundance-based estimator Chao1, Coverage: Good's coverage. lci and hci: lower and higher confidence interval, respectively.

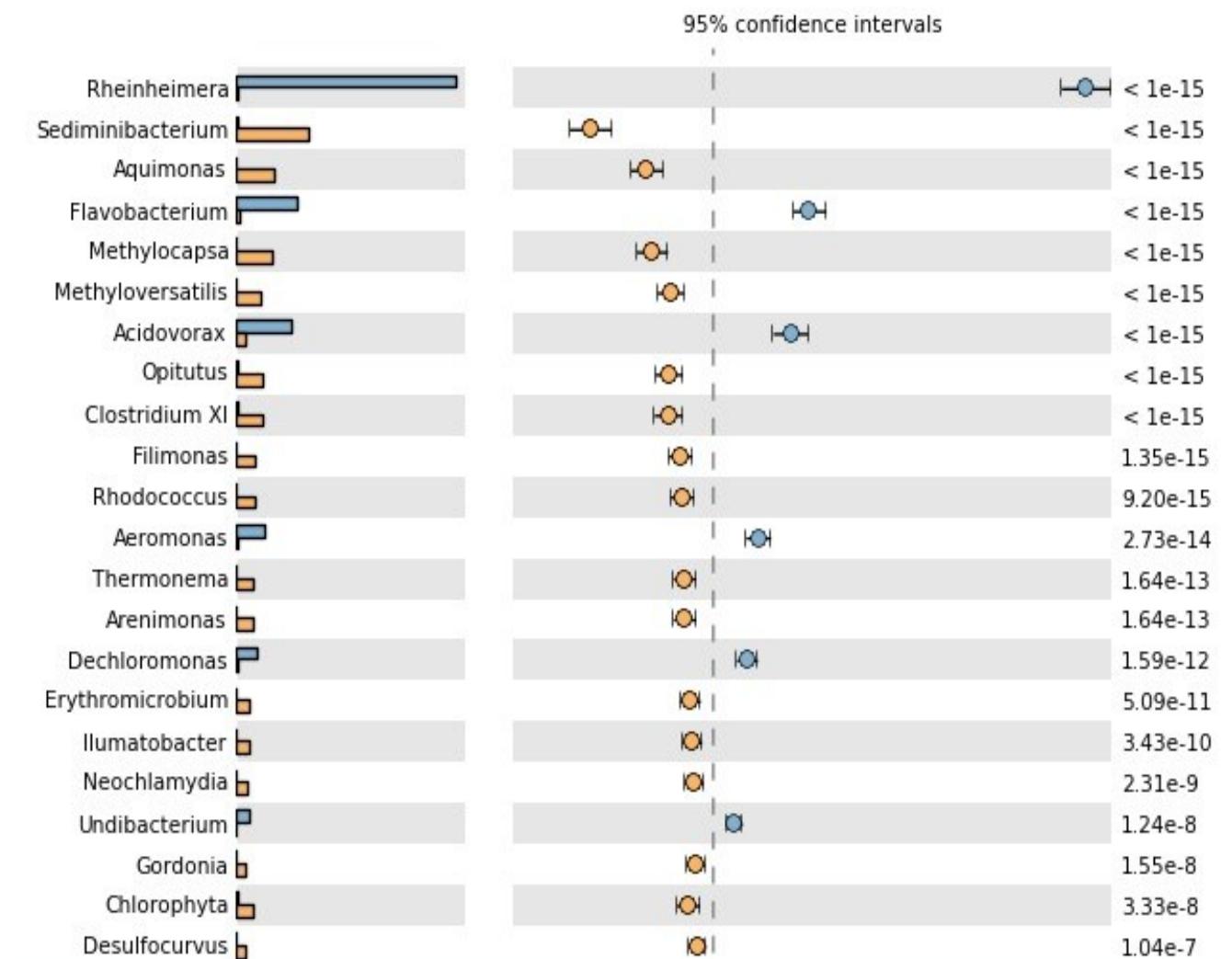
**Table S4** The 16 genera frequently identified in all biofilm samples from stone (ST) and tire derived rubber (TDR) media

Genus	Original Abundance	
	Stone (ST) biofilm	Tyre derived rubber (TDR) biofilm
<i>Reheinmera</i>	1019	99
<i>Rhodococcus</i>	26	840
<i>Aquabacterium</i>	4	830
<i>Trichococcus</i>	130	31
<i>Acidovorax</i>	296	167
<i>Flavobacterium</i>	242	43
<i>Aeromonas</i>	235	29
<i>Sediminibacterium</i>	143	52
<i>Hydrogenophaga</i>	94	123
<i>Aquimonas</i>	48	293
<i>Brevundimonas</i>	96	40
<i>Pseudoxanthomonas</i>	89	97
<i>Rhizobacter</i>	22	183
<i>Zoogloea</i>	19	13
<i>Arenimonas</i>	36	117
<i>Dechloromonas</i>	38	19
<i>Stenotrophomonas</i>	62	99

**Table S5** *P*-values between biofilm samples developed on tyre derived rubber (TDR) media at 10 and 30°C



**Key:** Blue color bars indicating *P*-values between biofilm samples developed at 10°C, Orange color bars indicating *P*-values between biofilm samples developed at 30°C.

**Table S6** *P*-values between biofilm samples developed on stone (ST) media at 10 and 30°C

**Key:** Blue color bars indicate *P*-values between biofilm samples developed at 10°C, Orange color bars indicate *P*-values between biofilm samples developed at 30°C.