1 AN OPEN AIR BIOCATHODE ENABLES EFFECTIVE ELECTRICITY

2 GENERATION WITH MICROBIAL FUEL CELLS

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Table S1: Overview of the organisms (O1-O9) corresponding with the sequenced bands on the DGGE represented in Figure S1.D. Identification was performed with the RDP Classification Algorithm. The probability values are given between brackets.

	Phylum	Order	Family	Genus
01	TM7 (100%)			Genera_incertae_sedis_TM7 (100%)
02	γ-Proteobacteria (100%)	Pseudomonadeles (100%)	Pseudomonadaceae (100%)	Pseudomonas (100%)
03	γ-Proteobacteria (100%)	Xanthomonadales (100%)	Xanthomonadaceae (100%)	Hydrocarboniphaga (100%)
04	α-Proteobacteria (100%)	Sphingomonoadeles (100%)	Sphingomonadaceae ((100%)	Novosphigobium (69%)
O 5	α-Proteobacteria (100%)	Sphingomonoadeles (100%)	Sphingomonadaceae (100%)	Sphingopyxis (100%)
06	α-Proteobacteria (100%)	Rhizobiales (100%)	Hyphomicrobiaceae (99%)	Xantobacter (95%)
07	α-Proteobacteria (98%)	Rhizobiales (83%)	Methylocystaceae (59%)	Albibacter (57%)
08	α-Proteobacteria (100%)	Rhodobacterales (100%)	Rhodobaceteraceae (100%)	Catellibacterium (54%)
09	α-Proteobacteria (100%)	Rhodobacterales (100%)	Rhodobaceteraceae (100%)	Catellibacterium (63%)

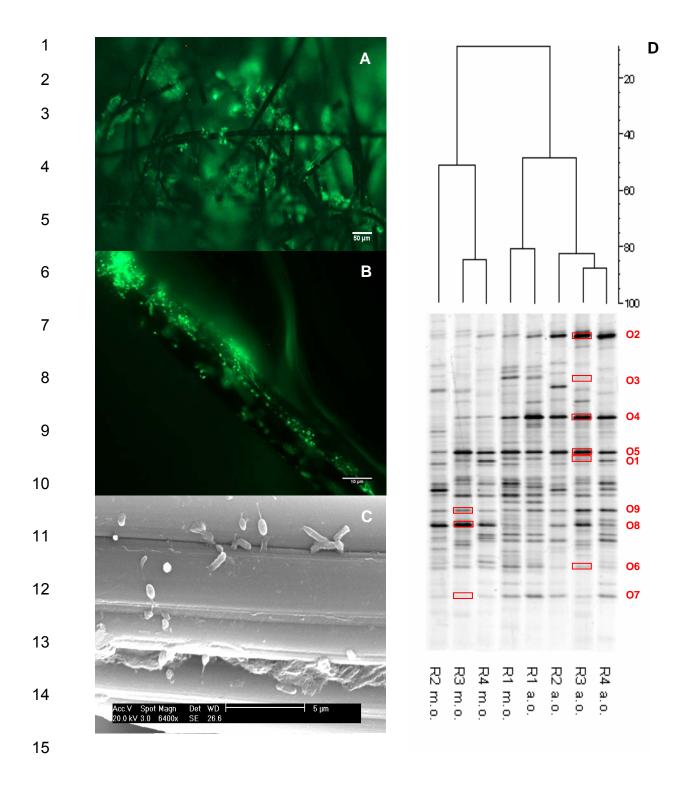


Figure S1: Epifluorescence microscopy images (A, B) and a secondary electron image (SEM) (C) of cathodic graphite fibers. DGGE pattern of the different biofilms on the cathodic felts after 5 months of operation; both the membrane oriented side (m.o.) as the air oriented side (a.o.) of the cathodic graphite felt were investigated (D) – The organisms (O1-O9) are indicated in Table S1

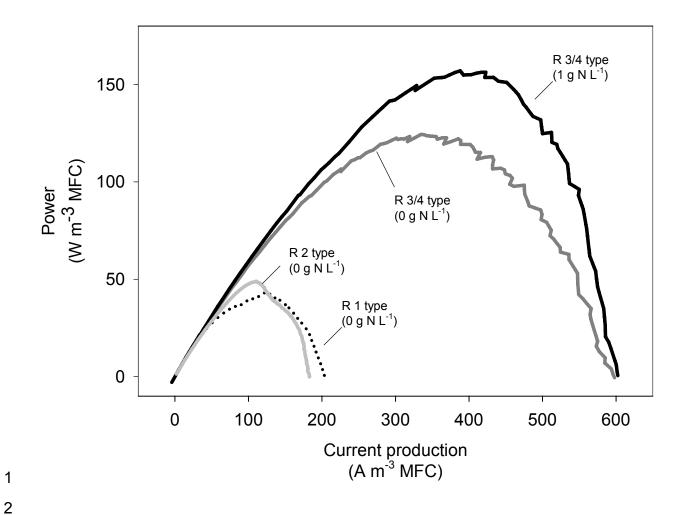


Figure S2: Power density curves (scan rate: 1 mV s⁻¹) for the different reactor designs (R1, R2 and R3/4) with 0.05 M ferricyanide as a cathodic electron acceptor (0.1 M phosphate buffer). The OCV was between 0.651 V and 0.708 V