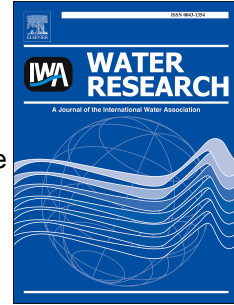


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New insights into the spatial variability of biofilm communities and potentially negative bacterial groups in hydraulic concrete structures

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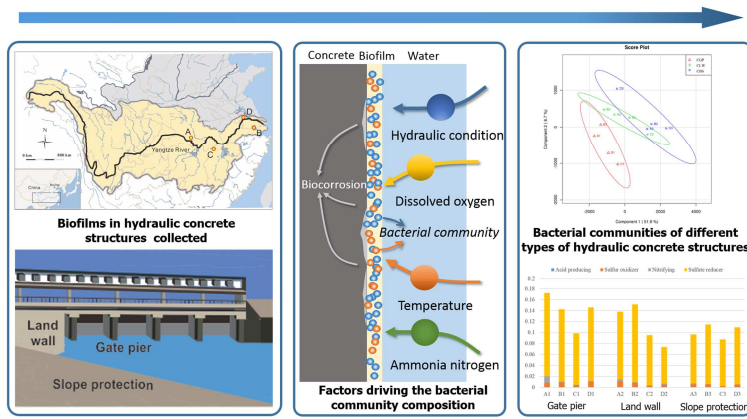
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