1 Supplementary Information:

Supplementary Fig. S1. DGGE analysis of AOB *amoA* gene fragments and their
transcripts from triplicate sediment samples collected from Hythe (Hy), Alresford (Al) and
Brightlingsea (Br) in the June (A), August (B) and October (C). Numbers indicate DGGE
bands that were excised and sequenced.

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Supplementary Fig. S2. DGGE analysis of AOB 16S rRNA gene fragments and their transcripts from triplicate sediment samples collected from Hythe (Hy), Alresford (AI) and Brightlingsea (Br) in the June (A), August (B) and October (C). Numbers indicate DGGE bands that were excised and sequenced.

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Supplementary Fig S3. A neighbor-joining tree of bacterial *amoA* (A) and 16S rRNA gene (B) sequences. The sequences retrieved during this study are shown in bold and include sample location, month. The letter/number codes are in accordance with the bands marked on the DGGE images in Supplementary Figs S1 and S2. Reference strains were obtained from public databases. The horizontal bar represents 5% sequence divergence. The bootstrapped consensus tree was inferred from 1000 replicates and the percentages over 50% are given at the nodes.

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Supplementary Fig. S4. Spatiotemporal patterns of diversity and composition in microbial communities along the Colne estuary. Seasonal changes in composition of bacteria (A) and archaea (C) communities are displayed on the ordination plots (visualised via an NMDS plot based on Jaccard's index). Symbols represent samples from July (circles), August (squares), October (triangles) and January (inverted triangles). Colours represent locations along the Colne estuary from the upper estuary (Hythe; black symbols) through the mid

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estuary (Alresford; grey symbols) to the estuary mouth (Brightlingsea; open symbols).
Seasonal changes in diversity of bacteria (B) and archaea (D) communities are shown
based on rarefied species richness estimates normalised to the sample with the fewest
amplicon reads, colours are the same as those in the ordination plots.



Supplementary Fig. S1



Supplementary Fig. S2



0.05

0.05

Supplementary Fig. S3

Α



