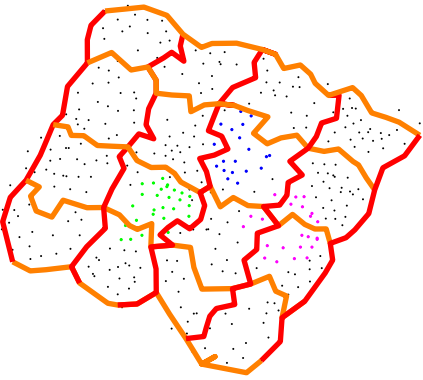
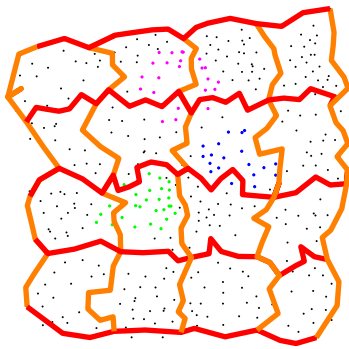


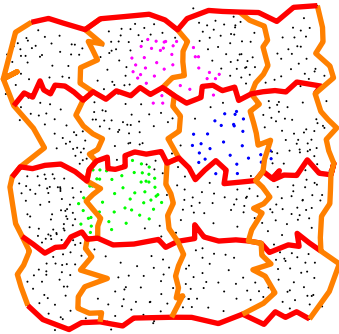
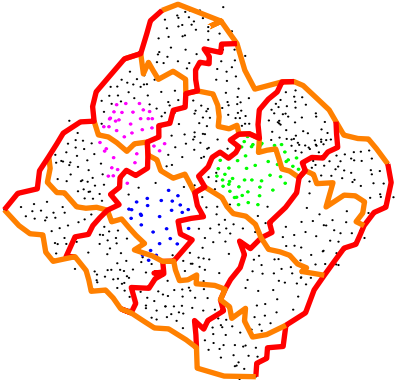
raw MDS output



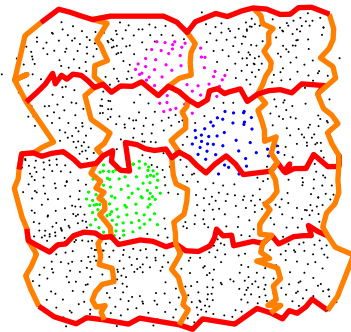
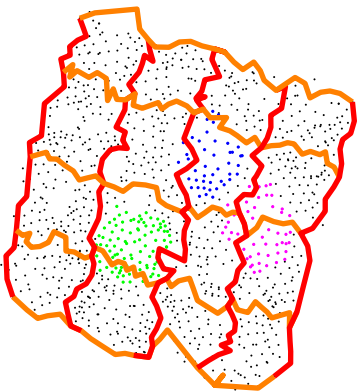
mismatch = 0.036, Ncells = 70



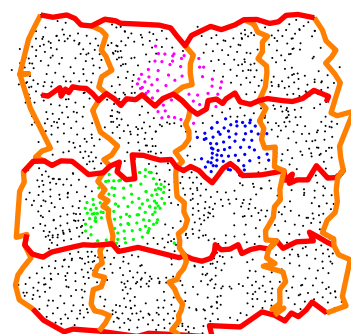
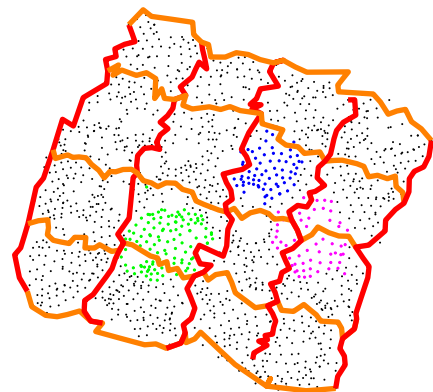
mismatch = 0.031, Ncells = 90



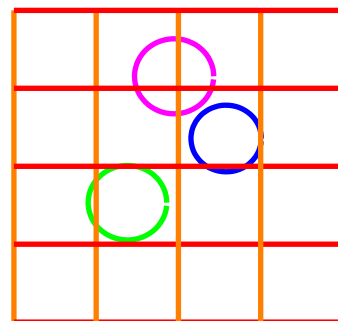
mismatch = 0.029, Ncells = 110



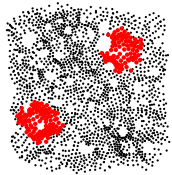
mismatch = 0.029, Ncells = 130



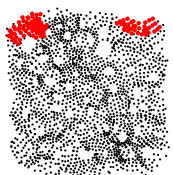
original space



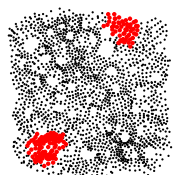
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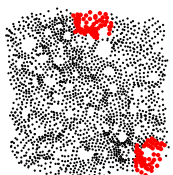
2



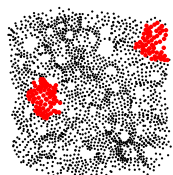
3



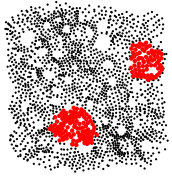
4



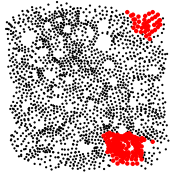
5



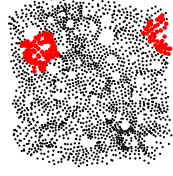
6



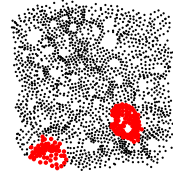
7



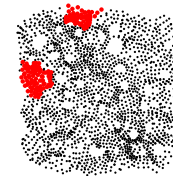
8



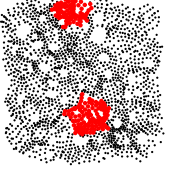
9



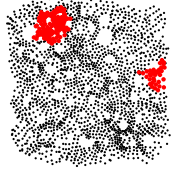
10



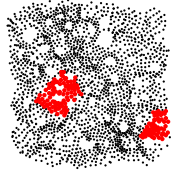
11



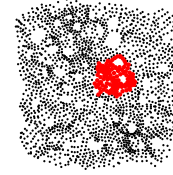
12



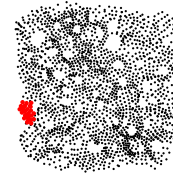
13

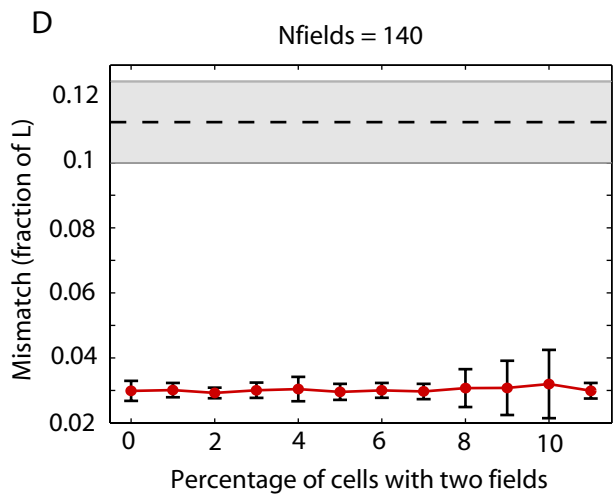
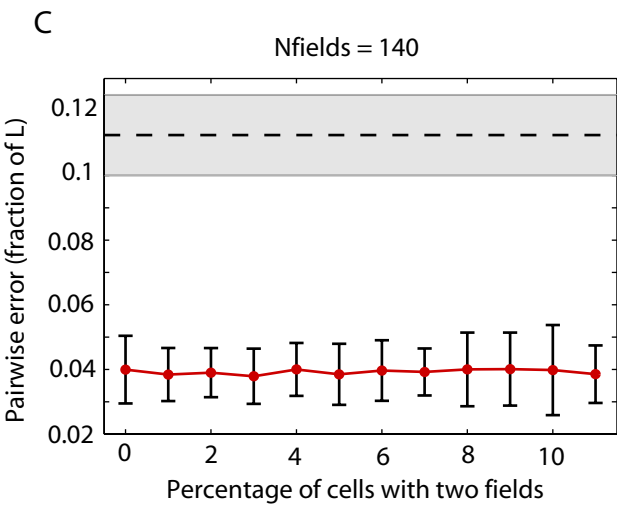
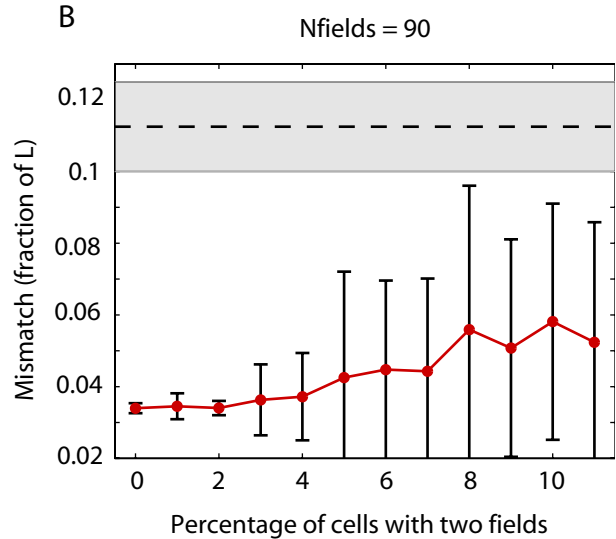
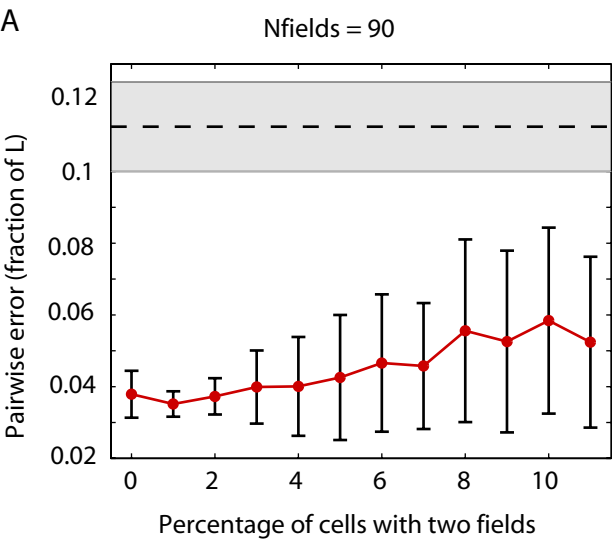


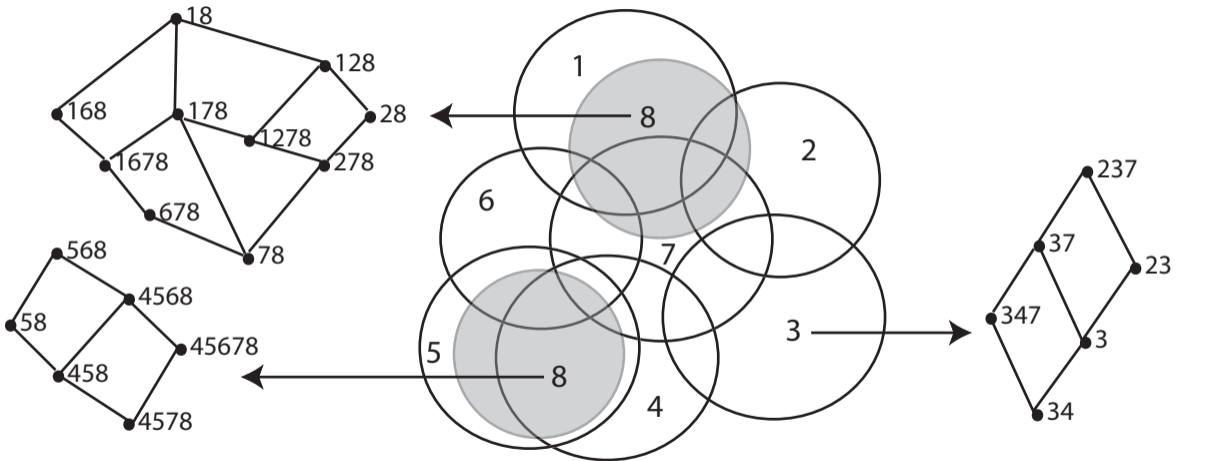
14



15







subgraph for cell 8

subgraph for cell 3