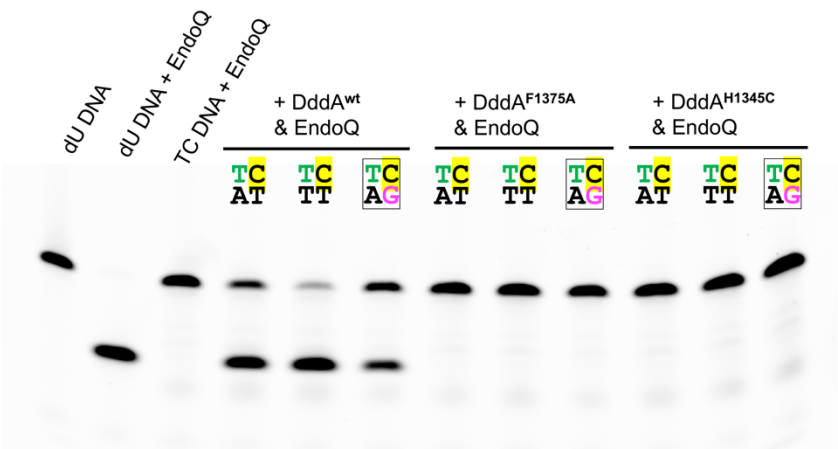
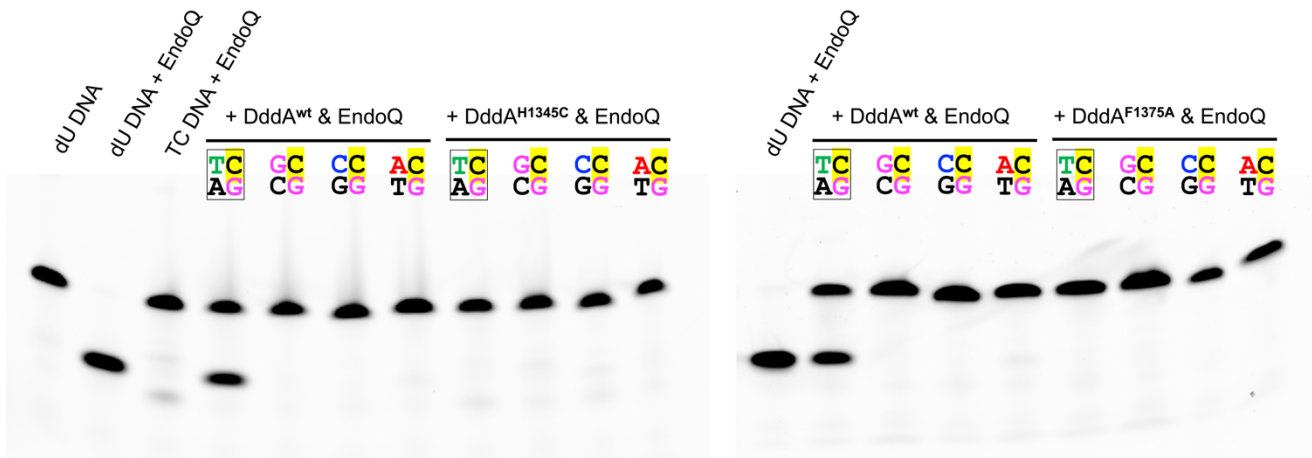




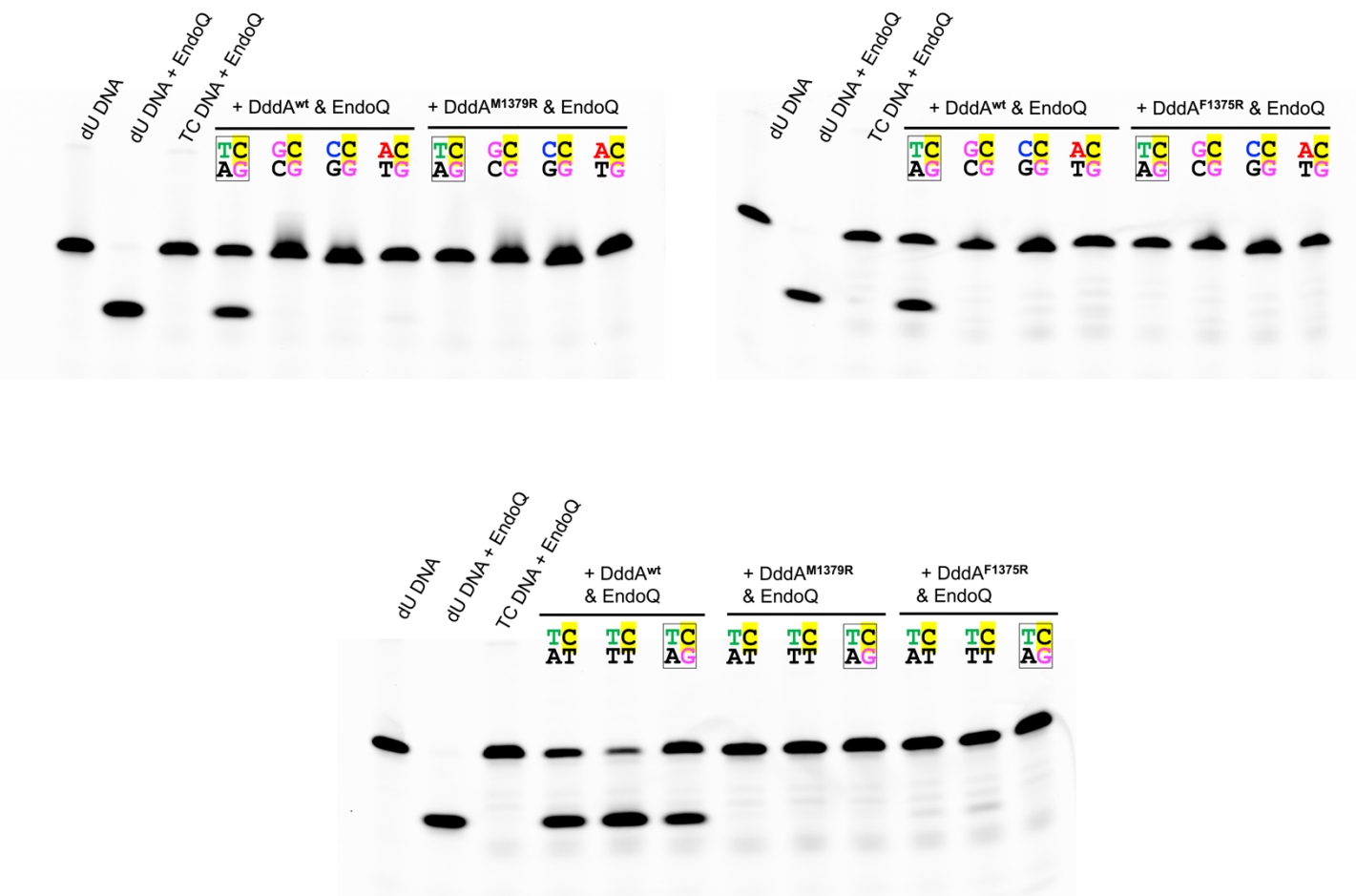
Structural basis of sequence-specific cytosine deamination by double-stranded DNA deaminase toxin DddA

In the format provided by the
authors and unedited

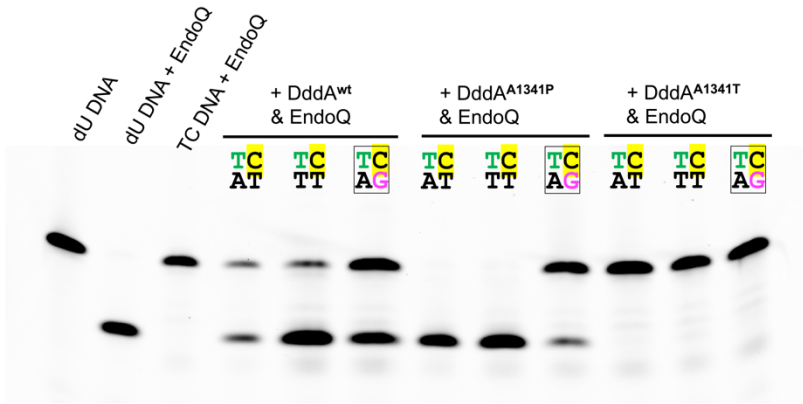
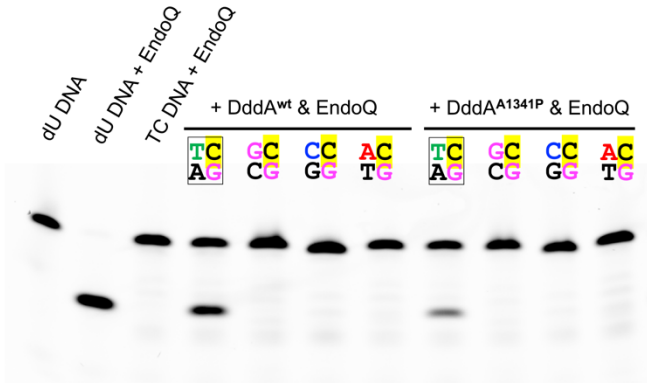
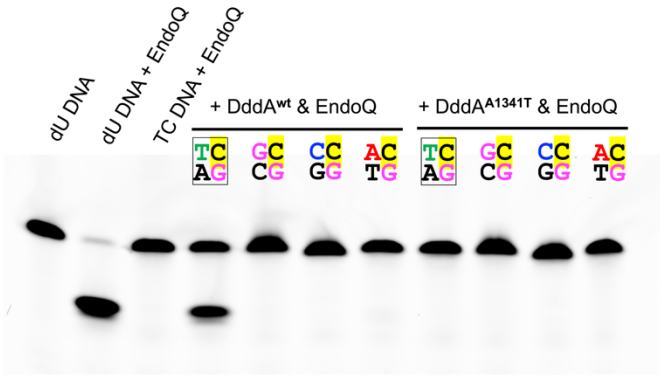
H1345C, F1375A



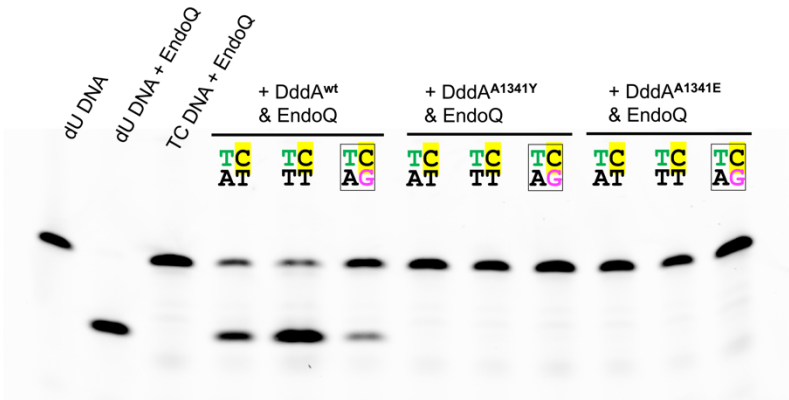
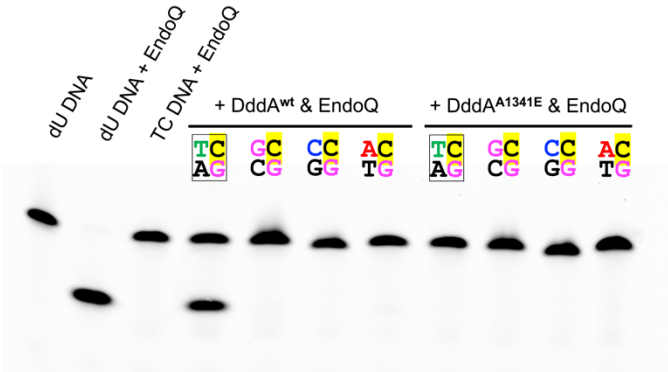
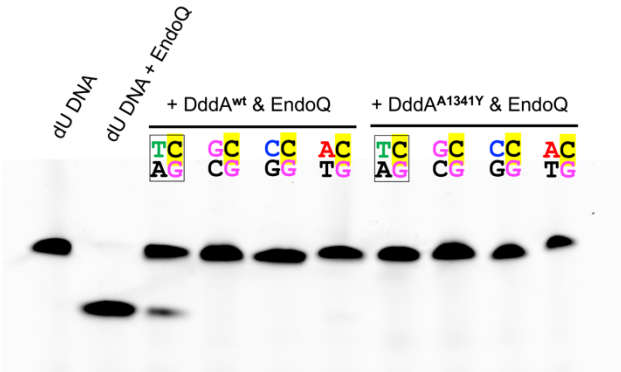
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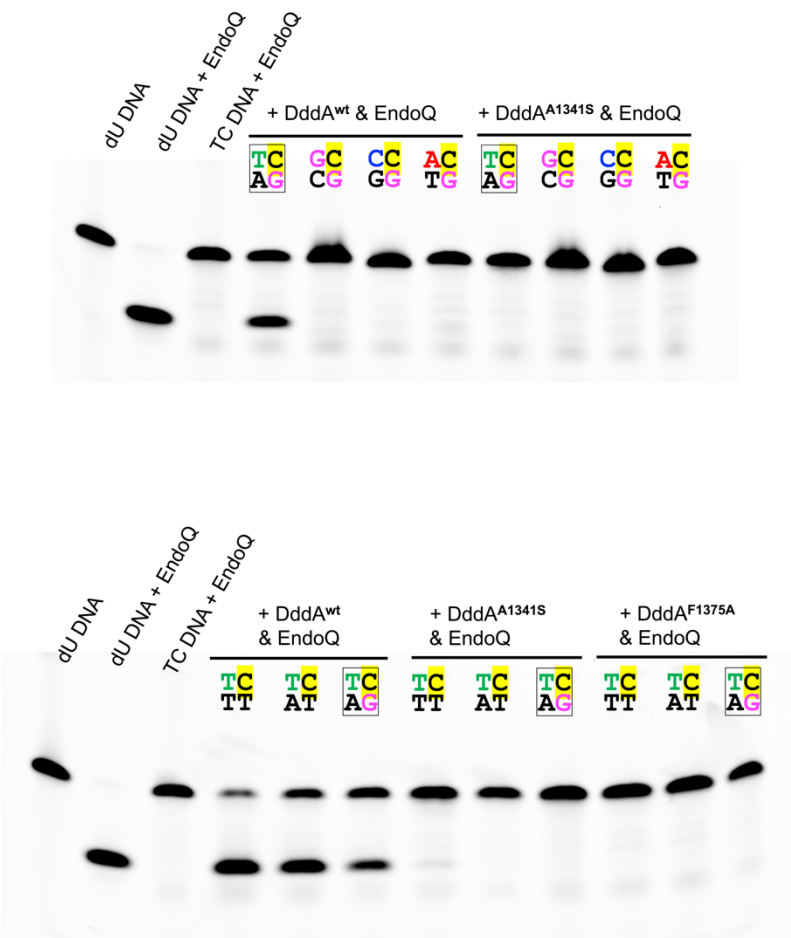
A1341T, A1341P



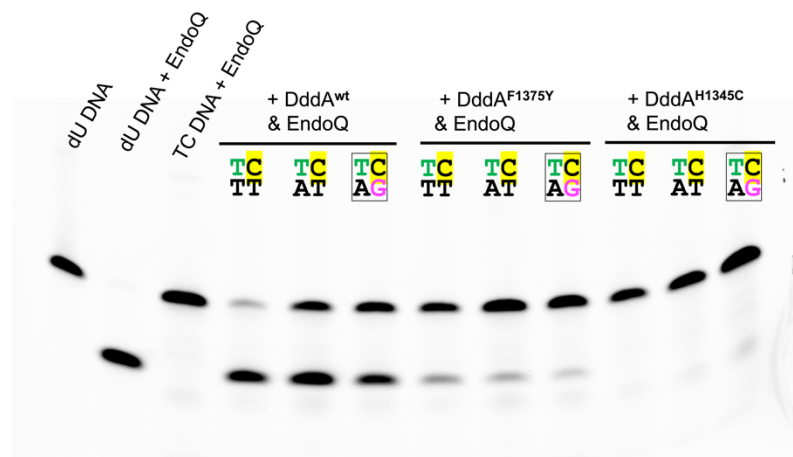
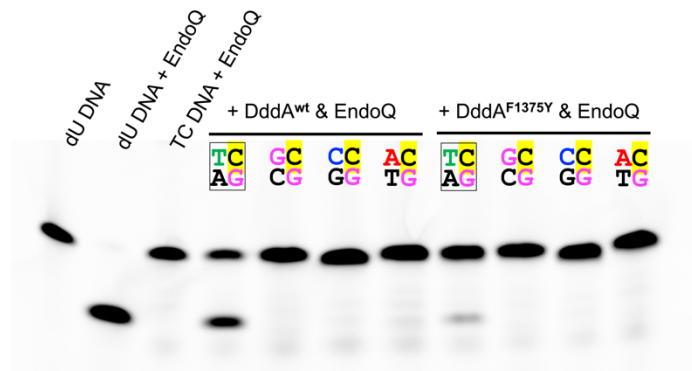
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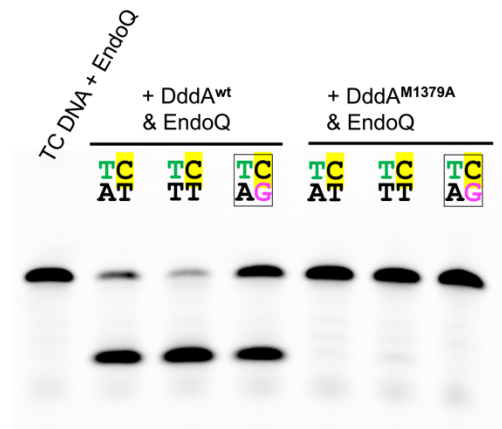
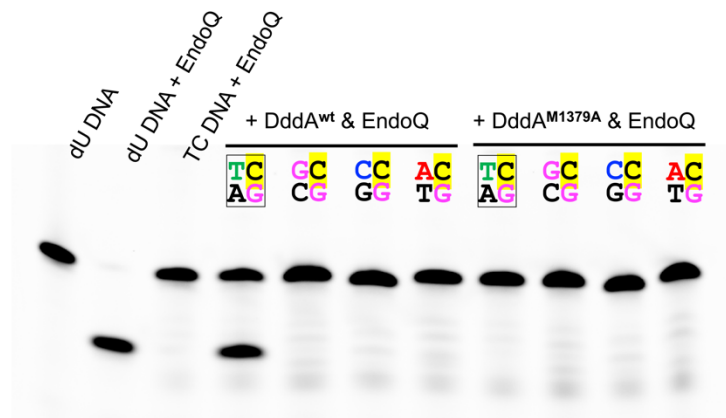
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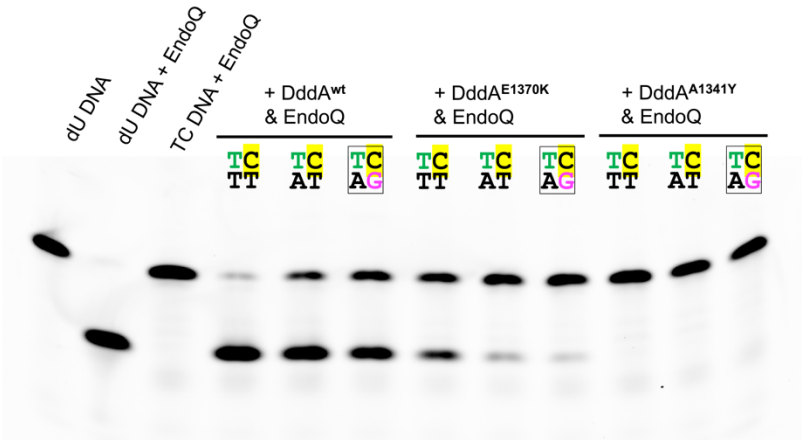
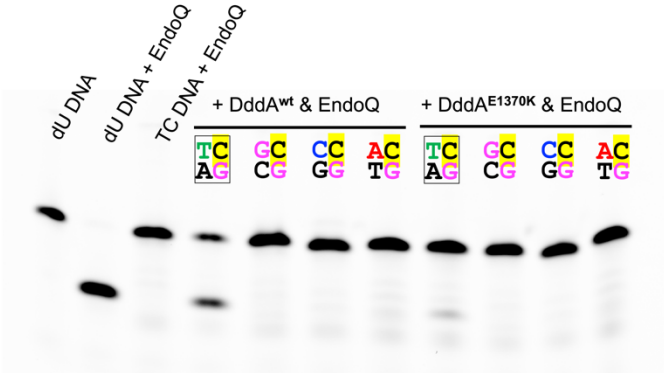
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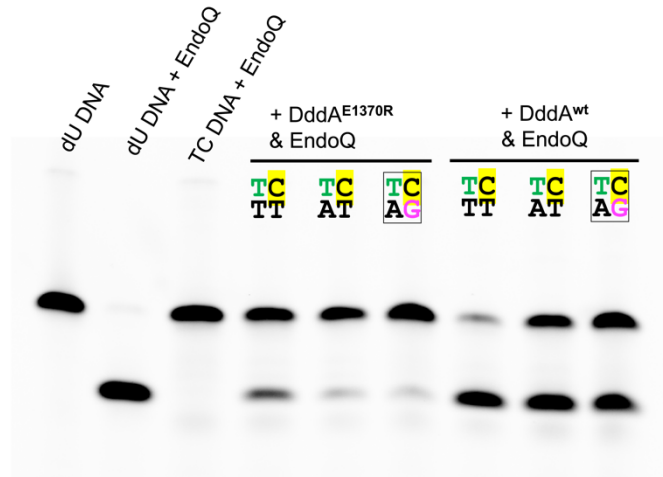
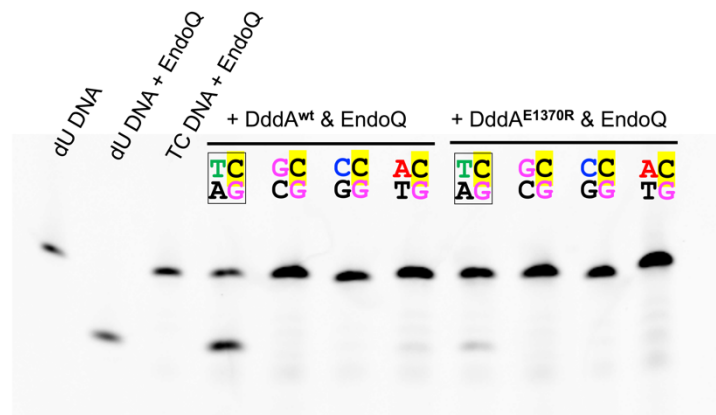
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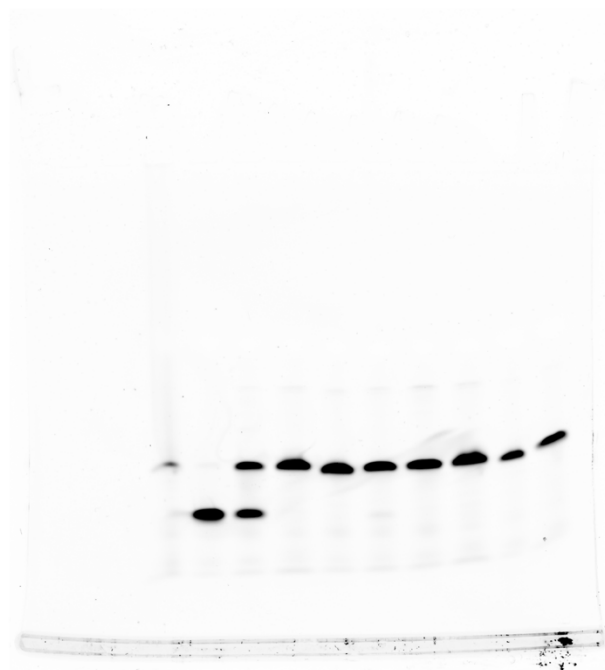


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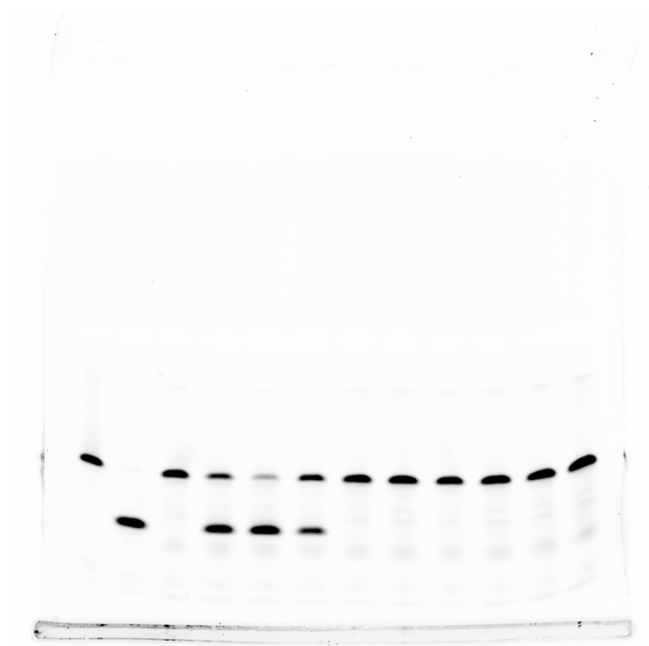


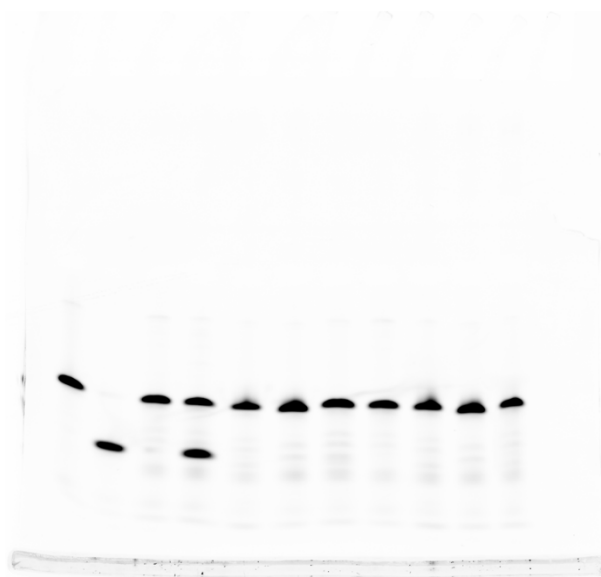
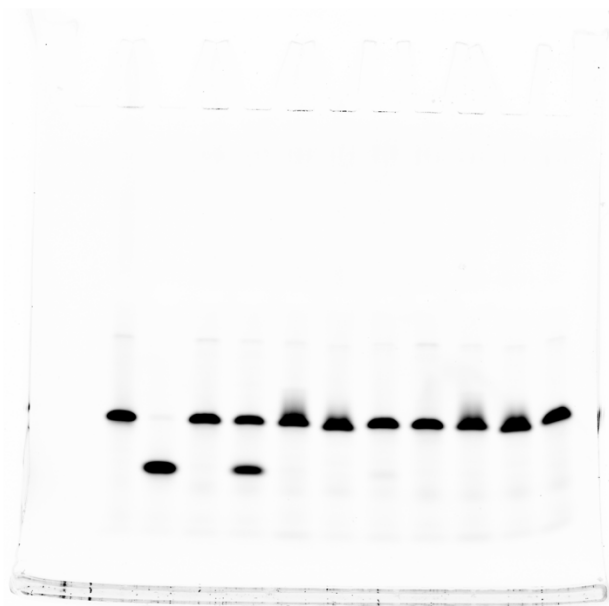
Supplementary Fig. 1 | Activities of various DddA mutants

The top and bottom strand sequences for the –1 and 0th positions are shown above each lane. Gels are representative of two replicates. Uncropped gel images are shown below.

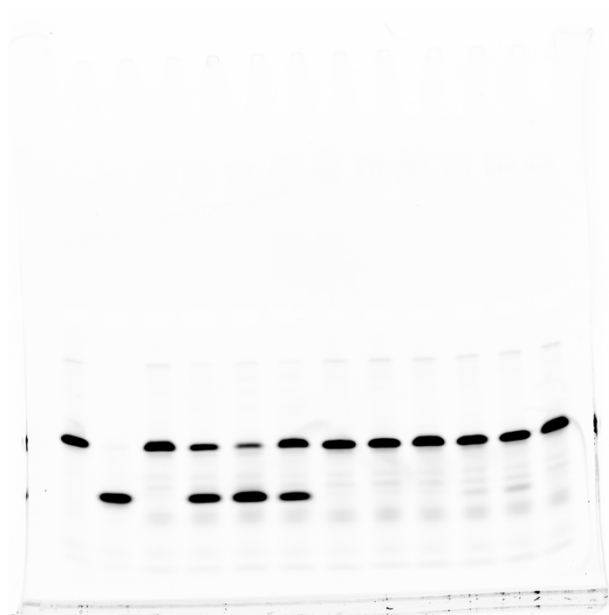


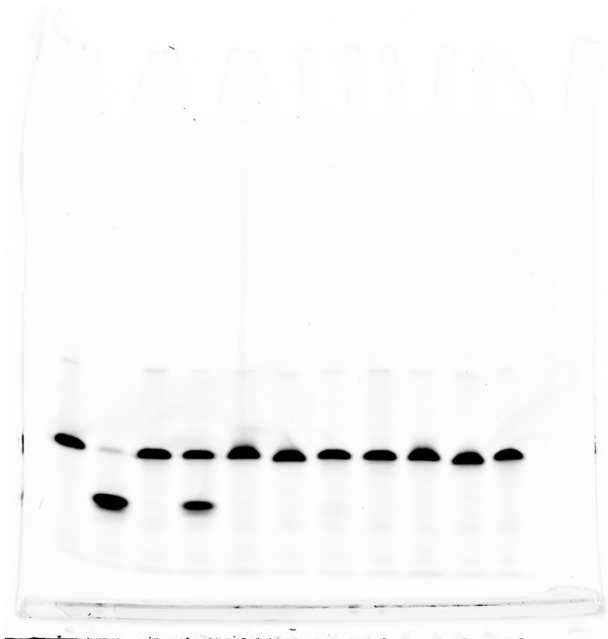
Uncropped gels
H1345C, F1375A





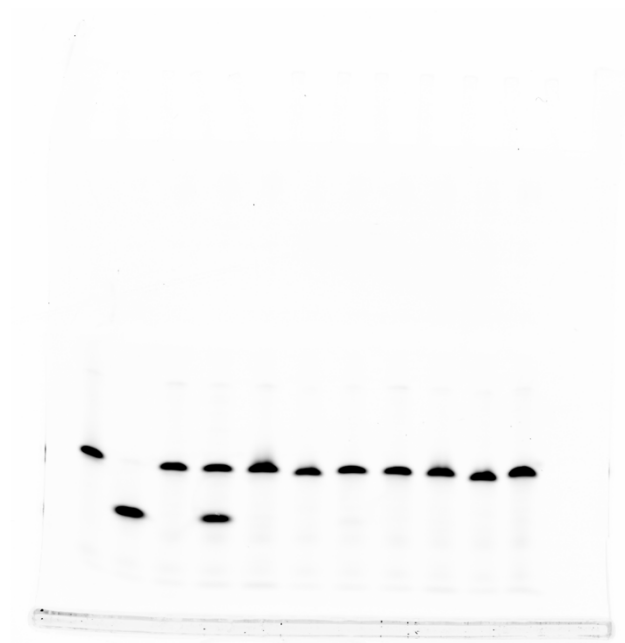
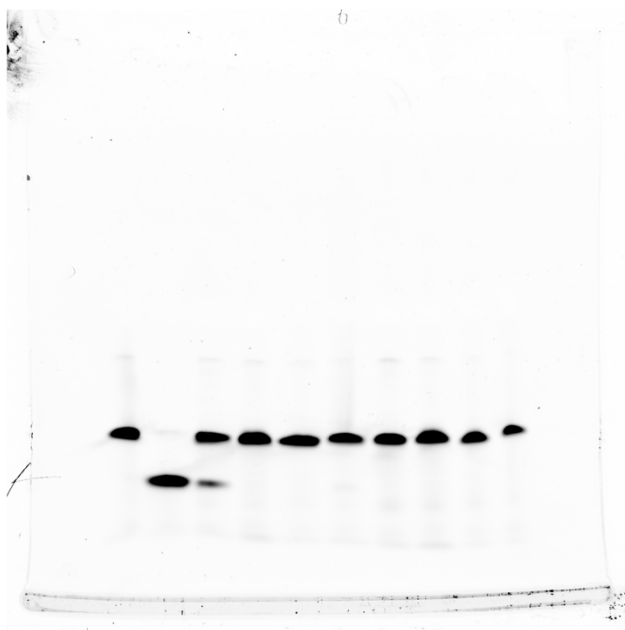
Uncropped gels
M1379R, F1375R



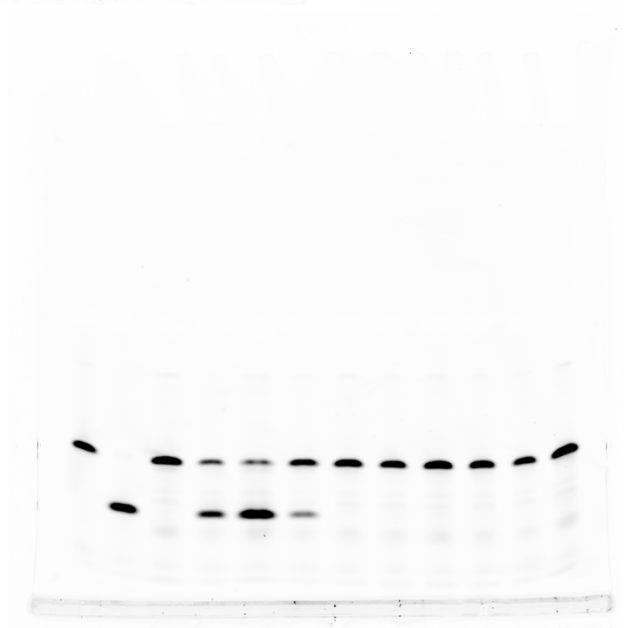


Uncropped gels
A1341T, A1341P

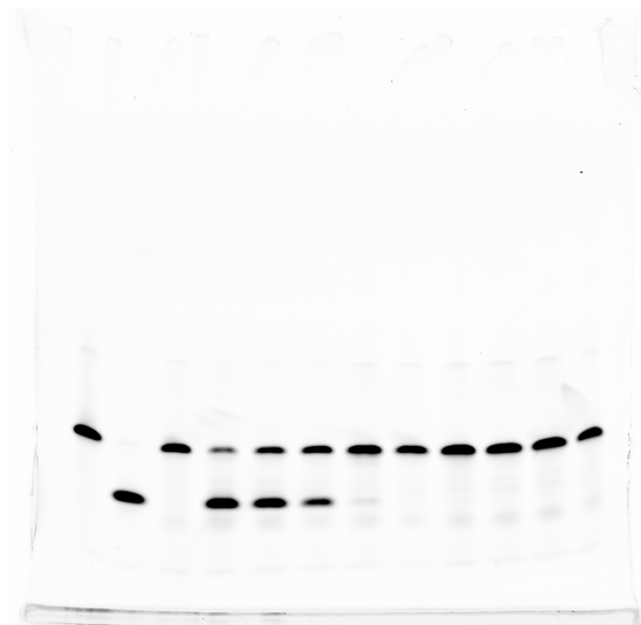
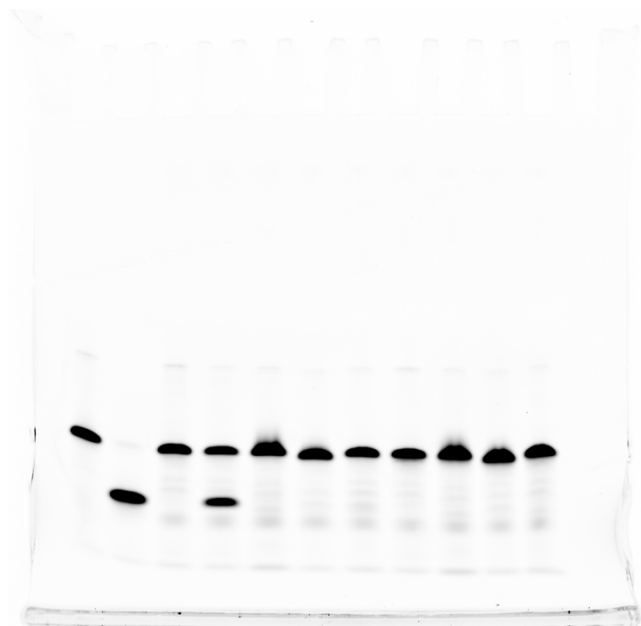




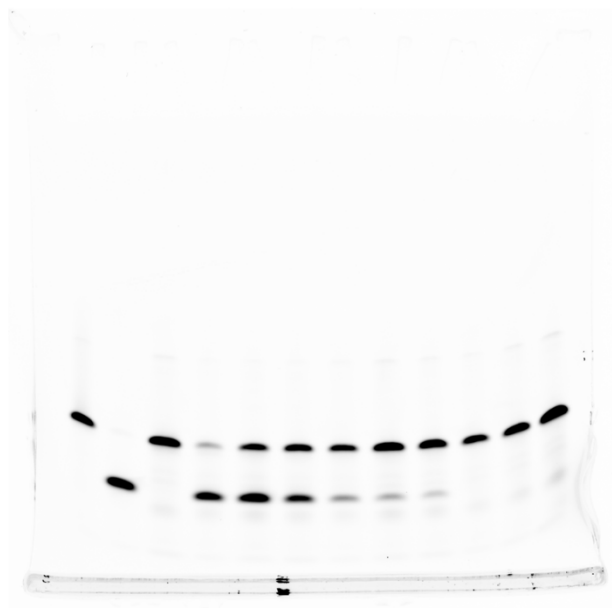
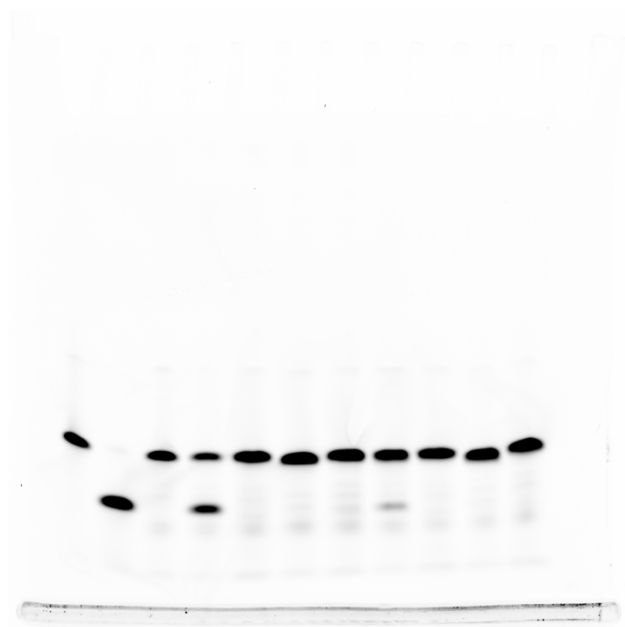
Uncropped gels
A1341Y, A1341E



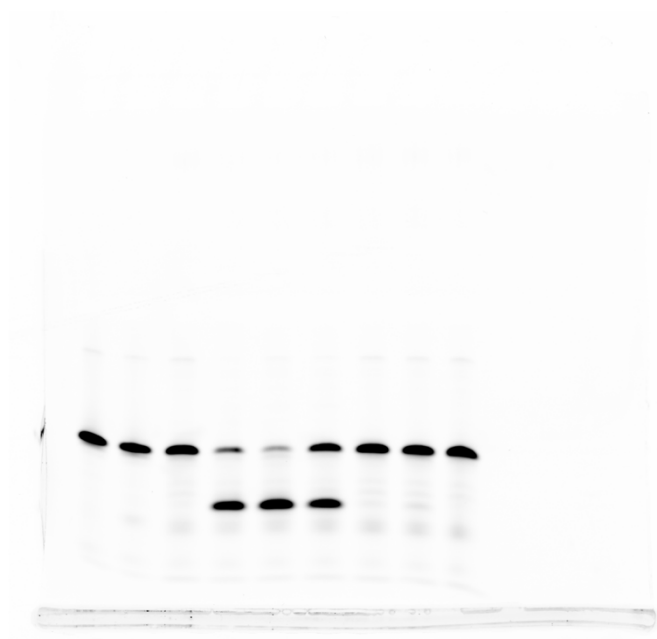
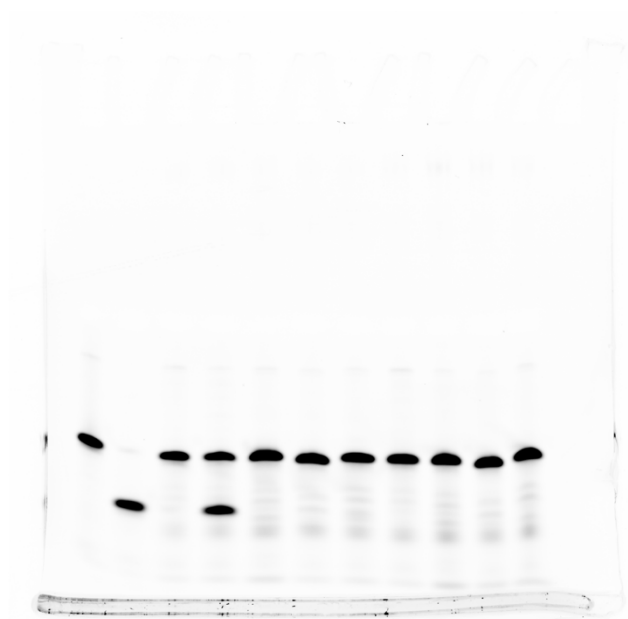
Uncropped gels
A1341S



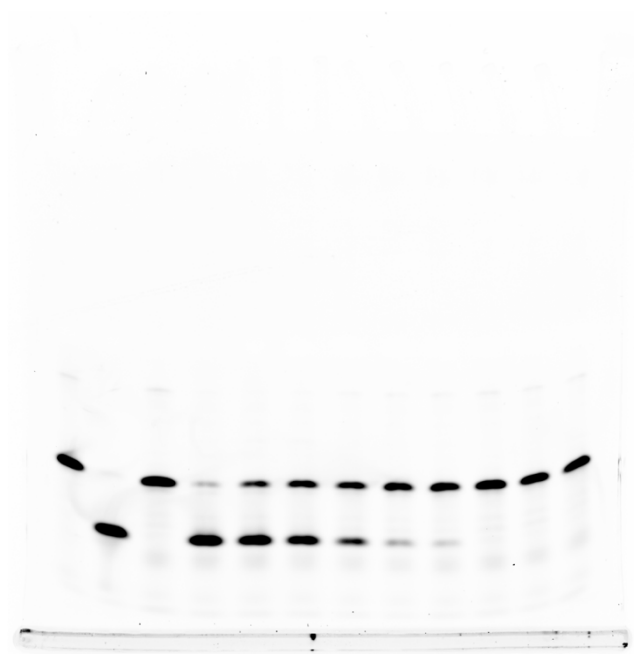
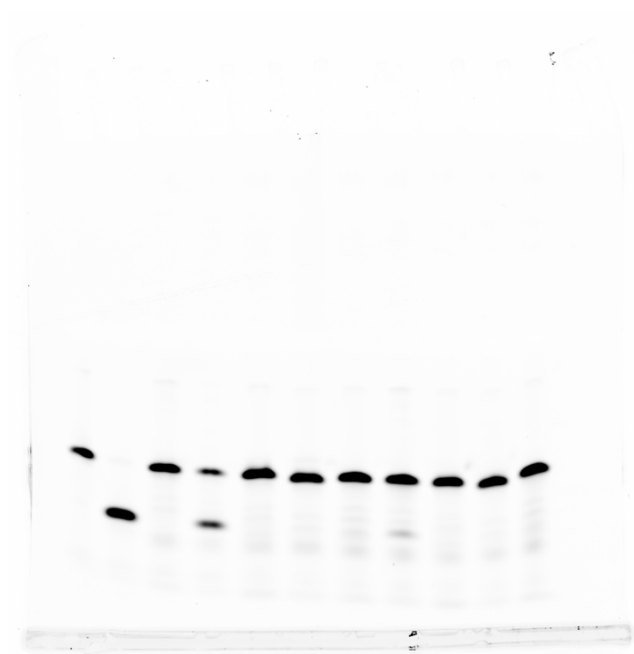
Uncropped gels
F1375Y



Uncropped gels
M1379A



Uncropped gels
E1370K



Uncropped gels
E1370R

