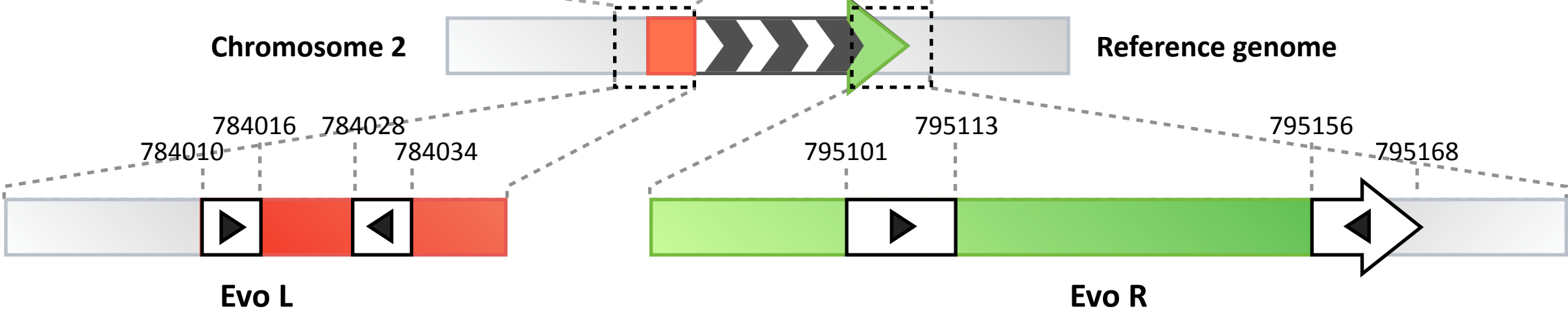


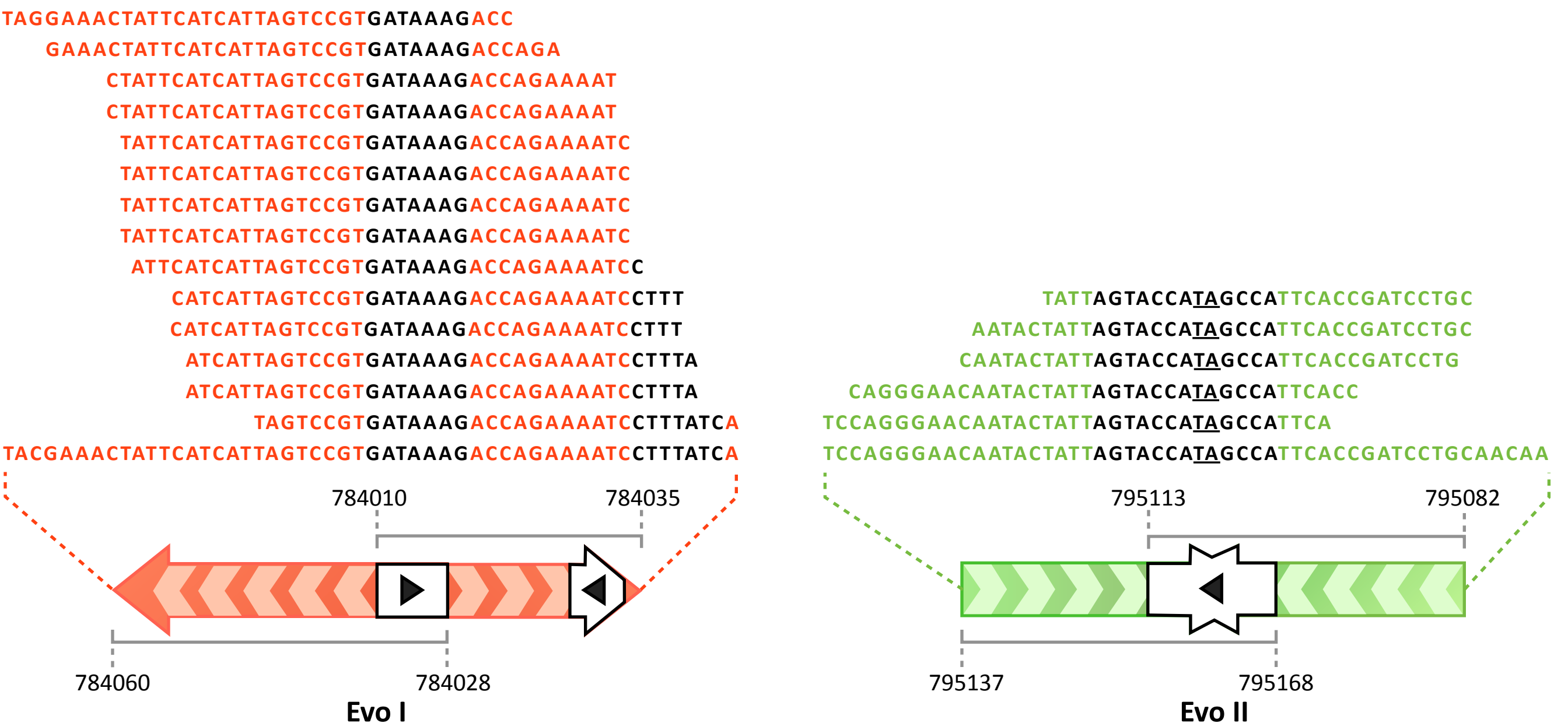
Sequence reads from evolved strain that match reference genome

ATTGTTGTTGCAGGATCGGTGAATGGCACTGGTACTTTGTTTGTGAAGCTGAGCAAGCTTCCAGGGAACAATACTATTAGTACCATAGCCACGATGGTTGATGAAGCA
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CTGAGCAAGCTTCCAGGGAACAATACTATTAGTACC
AGCAAGCTTCCAGGGAACAATACTATTAGTACCATA
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CCAGGGAACAATACTATTAGTACCATAGCCACGATG
GAACAATACTATTAGTACCATAGCCACGATGGTTGA
ACTATTAGTACCATAGCCACGATGGTTGATGAAGCA

CATGGAACTGGCAAAACACCACGATAAAGACCAGA
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CACGATAAAGACCAGAAAATCCTTTATCACGGACTA
GATAAAGACCAGAAAATCCTTTATCACGGACTAATG
CATGGAACTGGCAAAACACCACGATAAAGACCAGAAAATCCTTTATCACGGACTAATG



Contigs of unmapped reads from evolved strain



Proposed structure of the rearrangement

