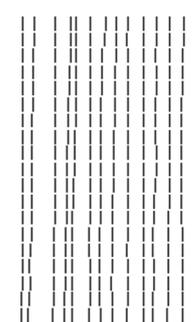


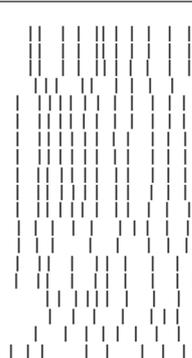
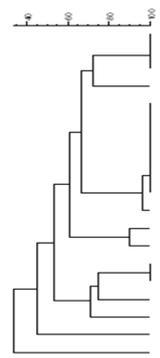
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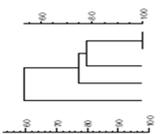
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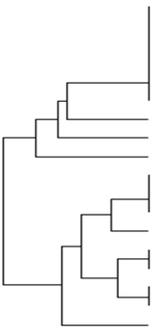
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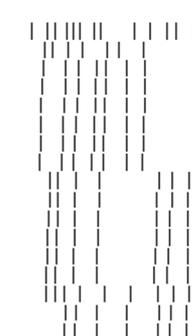
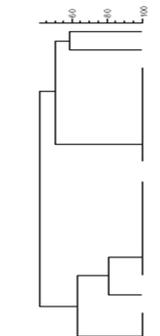
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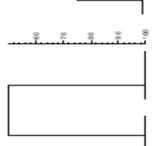
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 150\_brine\_Y28.5  
 150\_brine\_Y28.6  
 150\_brine\_Y28.7  
 starter\_Y28\_30\_07u  
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 150\_brine\_Y28.15  
 150\_brine\_Y28.16  
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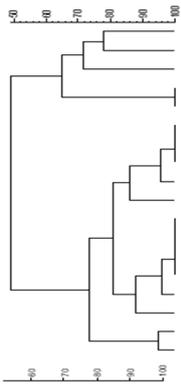
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 75\_brine\_Y28.4  
 75\_brine\_Y28.5  
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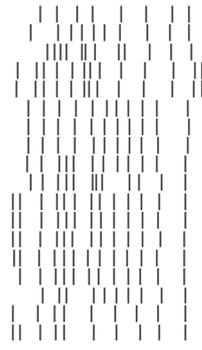
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(B)

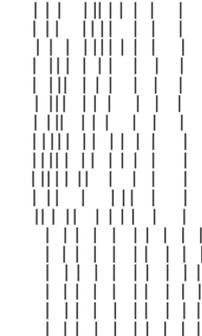
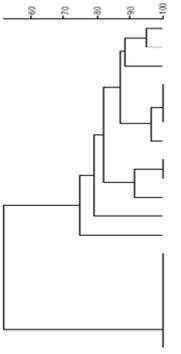
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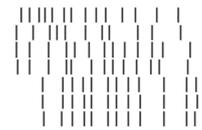
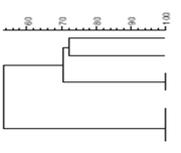
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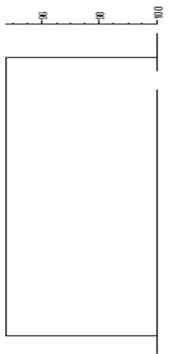
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 150\_olive\_Y30.2rep



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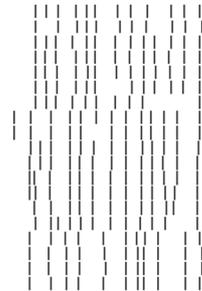
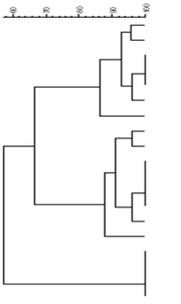
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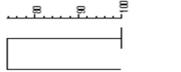
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 0\_Brine\_Y30.7  
 0\_Brine\_Y30.8  
 0\_Brine\_Y30.9  
 starter\_Y30\_10.07



0\_Brine\_Y30.18  
 0\_Brine\_Y30.19  
 0\_Brine\_Y30.20  
 starter\_Y30A\_10.07m



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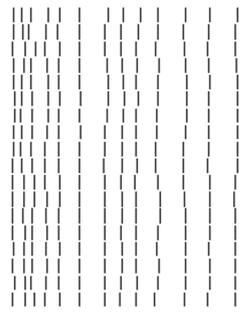
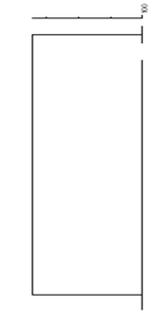


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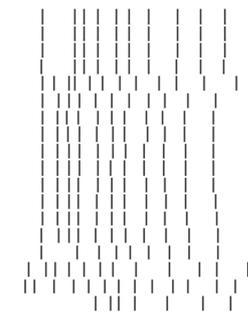
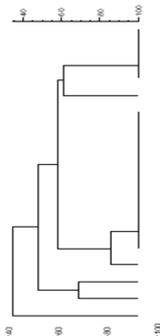
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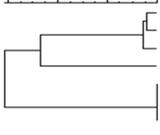
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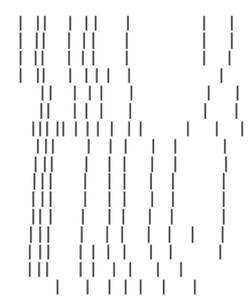
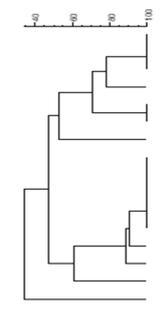
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 0\_brine\_Y31.4  
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 starter\_Y31A\_13.07u



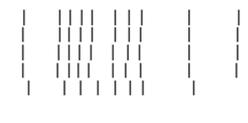
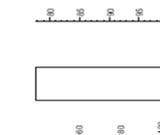
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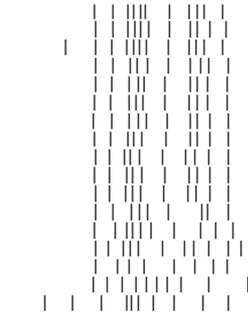
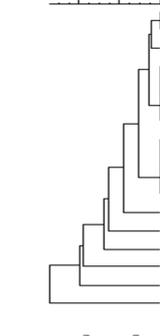
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 starter\_Y31\_13.07d



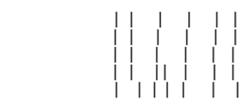
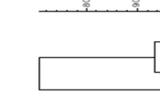
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 starter\_Y31\_20.07m  
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 75\_brine\_Y31.13  
 75\_brine\_Y31.16



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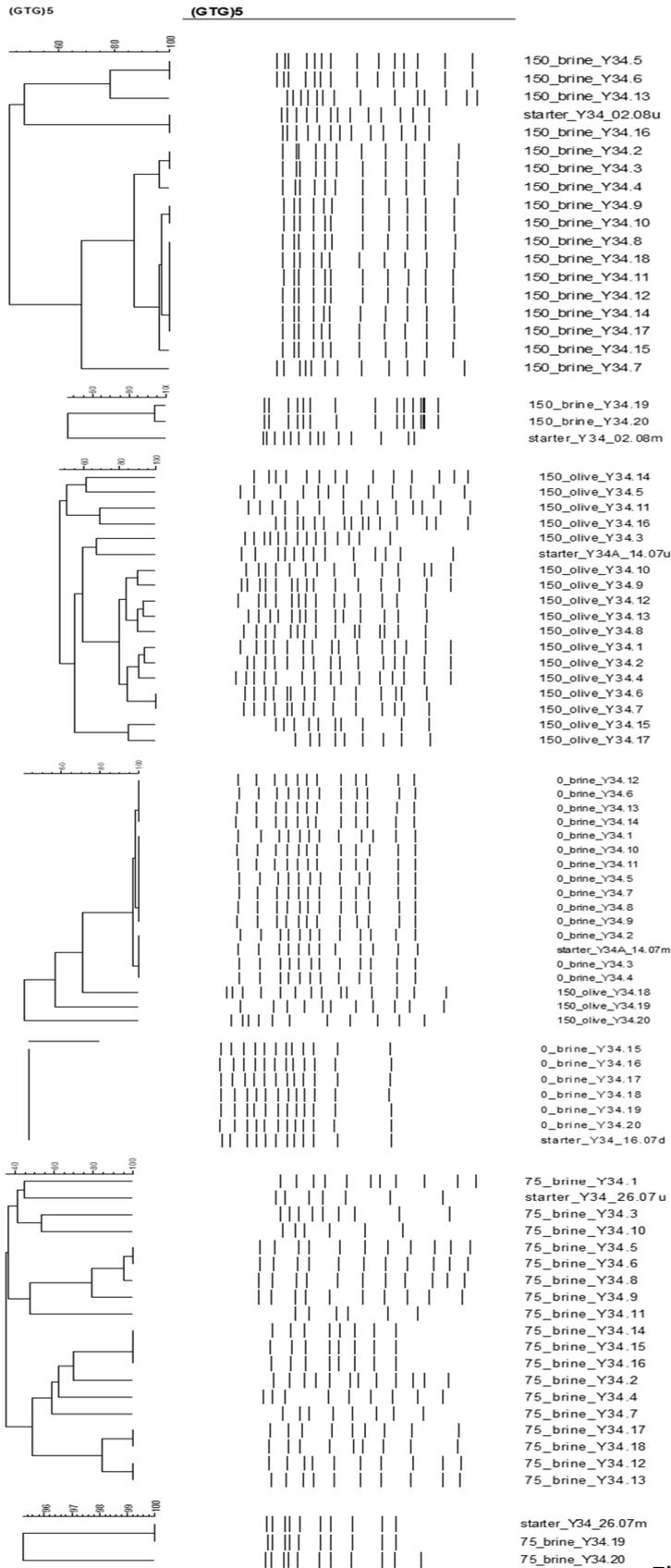


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 150\_brine\_Y31.14  
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 150\_brine\_Y31.15  
 starter\_Y31\_30.07Bm



150\_brine\_Y31.17  
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 150\_brine\_Y31.20  
 150\_brine\_Y31.19  
 starter\_Y31\_30.07Bm

(D)



**Figure S1.** Cluster analysis of rep-PCR patterns of the isolates comparing to starters for the three sampling points (0, 75, and 150 days) in the brines and olives for the five inoculated fermentations: (A) *C. boidinii* Y27, (B) *C. boidinii* Y28, (C) *C. boidinii* Y30, (D) *C. boidinii* Y31, and (E) *S. cerevisiae* Y34. Distance is indicated by the mean correlation coefficient [r (%)] and clustering is performed by UPGMA analysis.