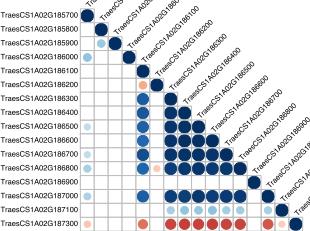
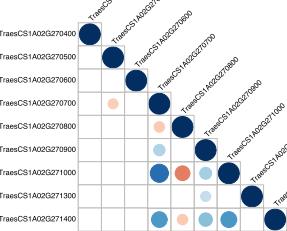


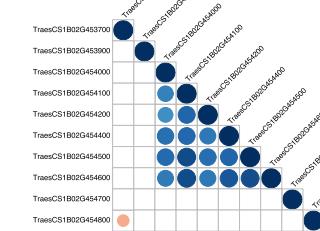
FRGC1



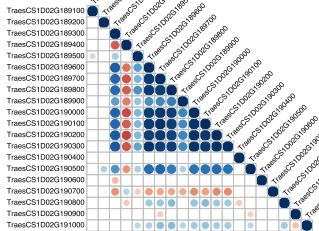
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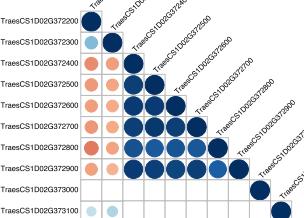
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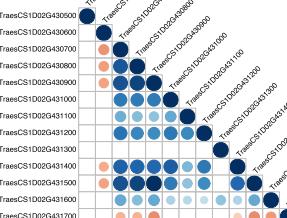
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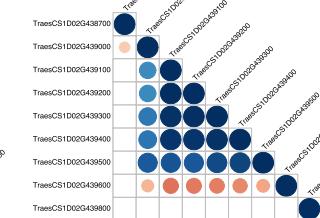
FRGC5



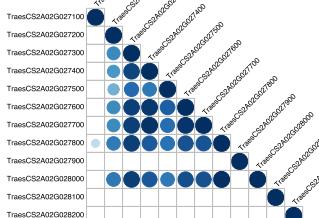
FRGC6



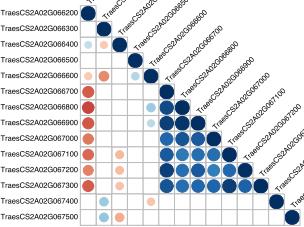
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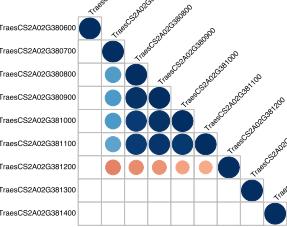
FRGC8



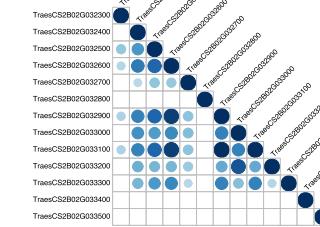
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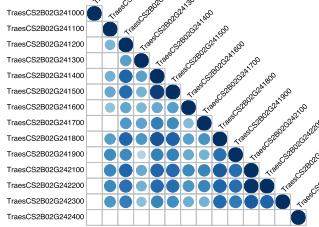
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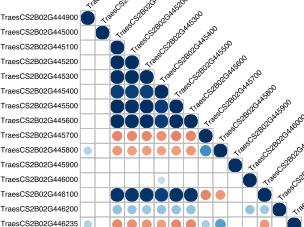
FRGC11



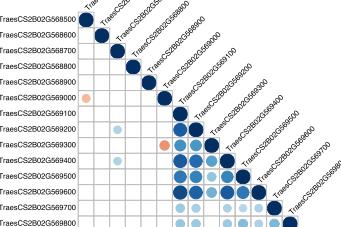
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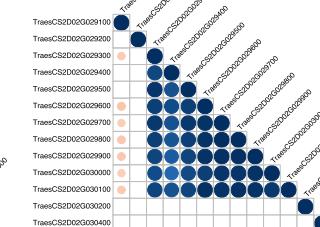
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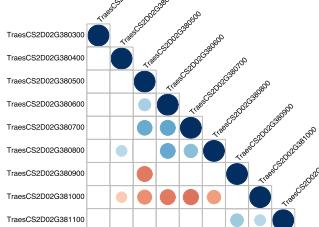
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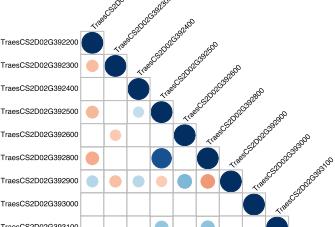
FRGC15



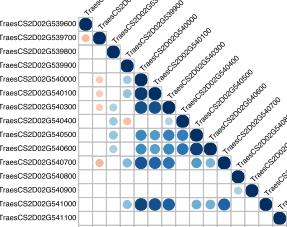
FRGC16



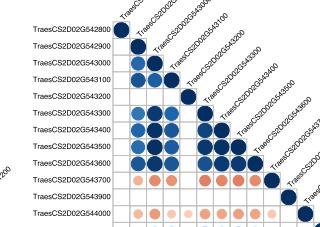
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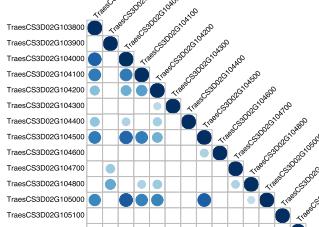
FRGC18



FRGC19

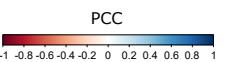
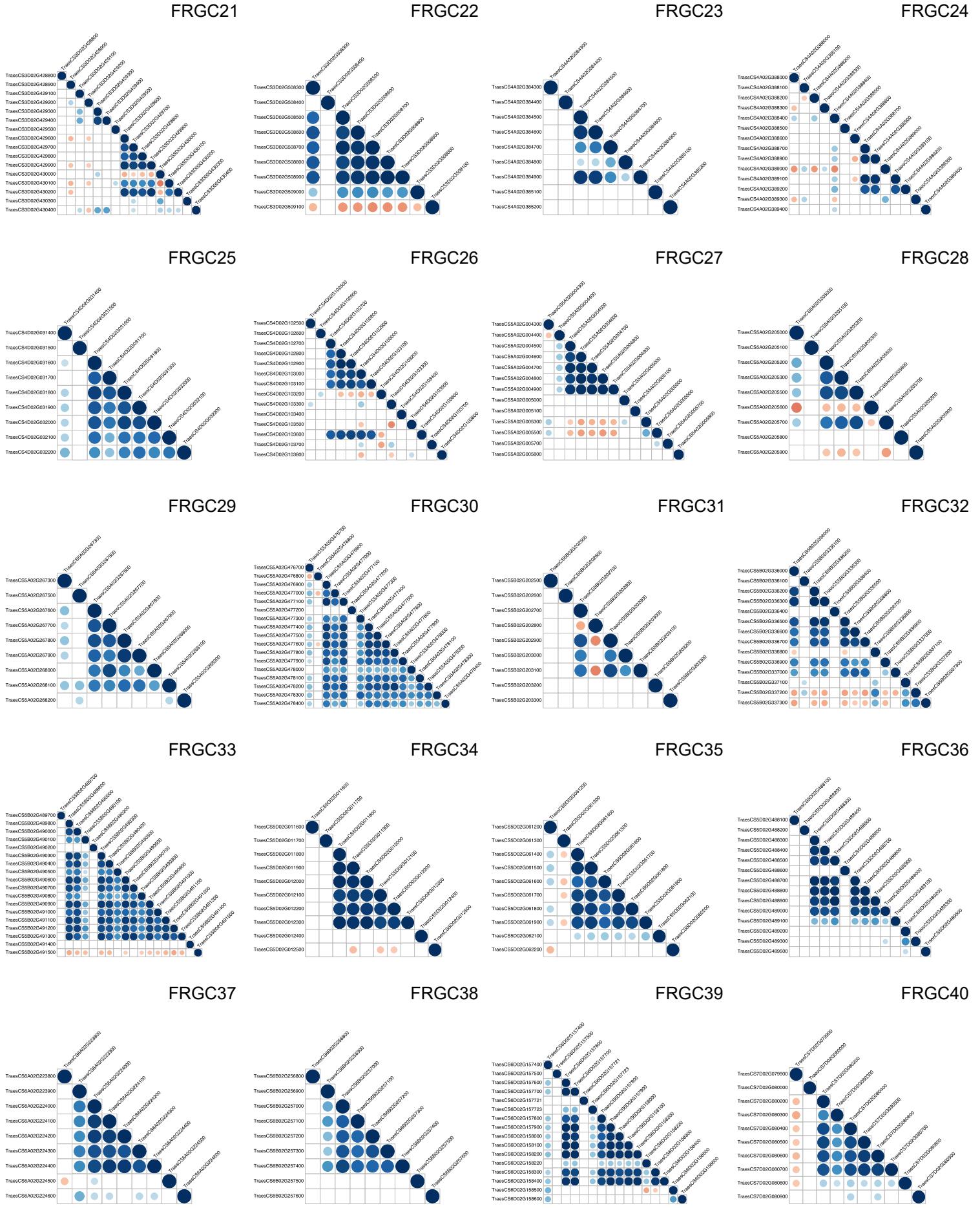


FRGC20

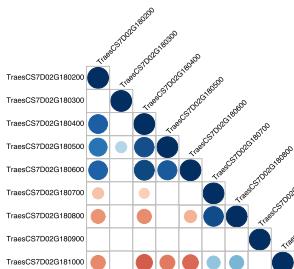


PCC

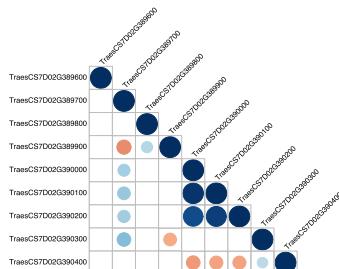
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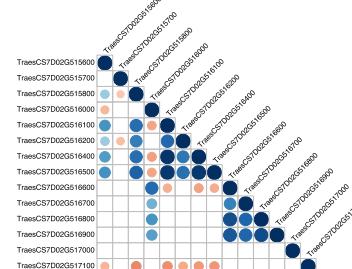
FRGC41



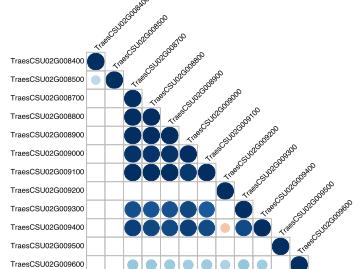
FRGC42



FRGC43



FRGC44



PCC

-1 -0.8 -0.6 -0.4 -0.2 0 0.2 0.4 0.6 0.8 1

Figure S2. Co-expression matrix of FRGC. Expression of clustered genes plus the two neighbouring genes flanking each side of the cluster were used to calculate to generate a co-expression matrix for each FRGC. Circles characteristics represent the direction (colour, blue = positive, red = negative) and the strength (size) of the correlation.