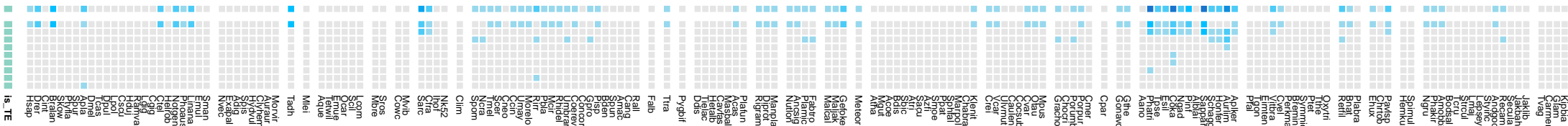
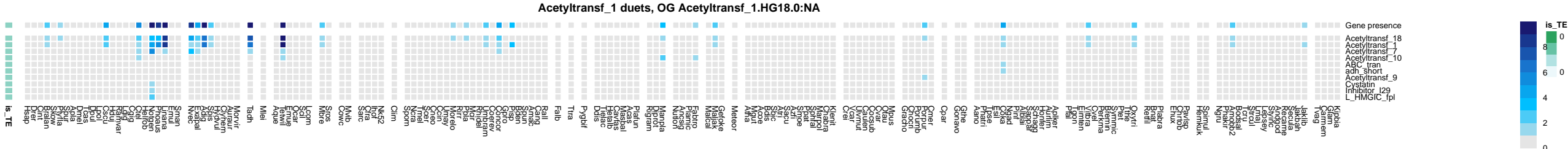
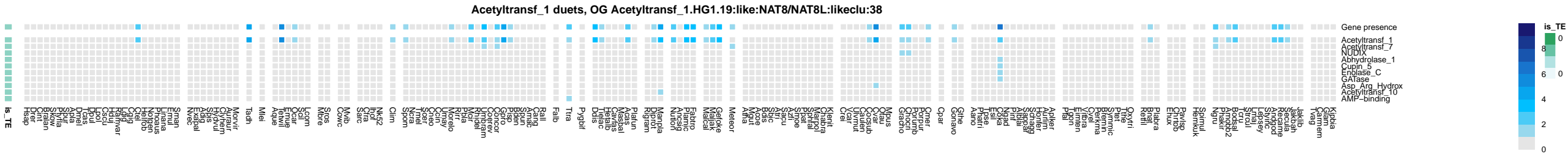


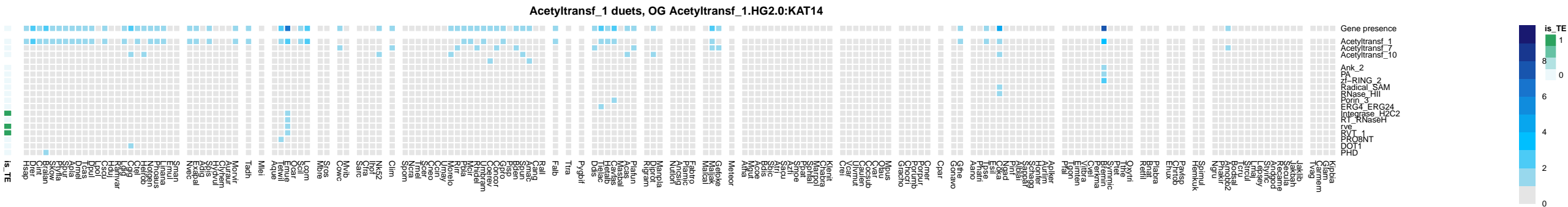
Gene presence

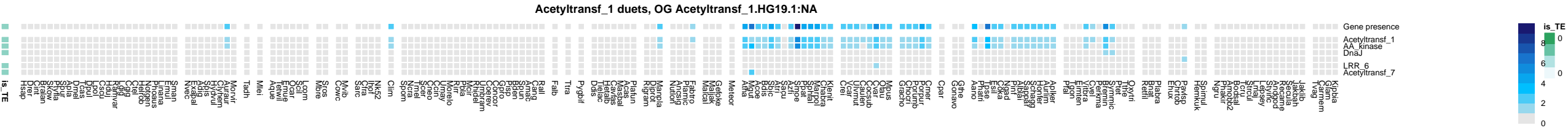
Acetyltransf\_1  
PhzC-PhzF  
Acetyltransf\_10  
Acetyltransf\_7  
Pirin  
Pirin\_C  
DfW  
RINGv  
Acetyltransf\_9

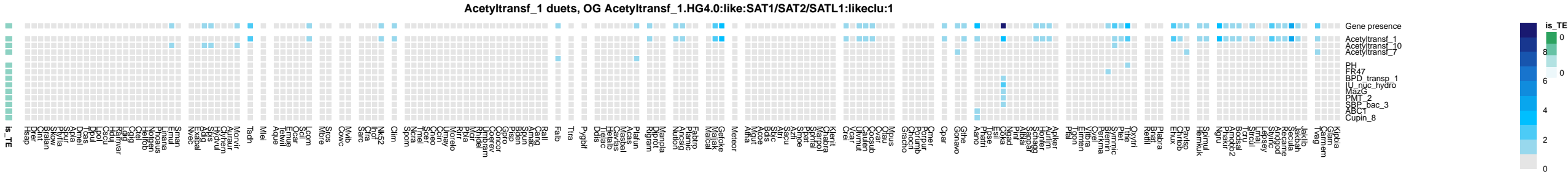


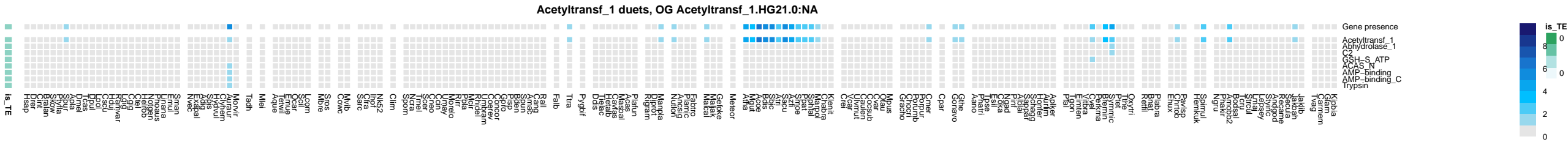


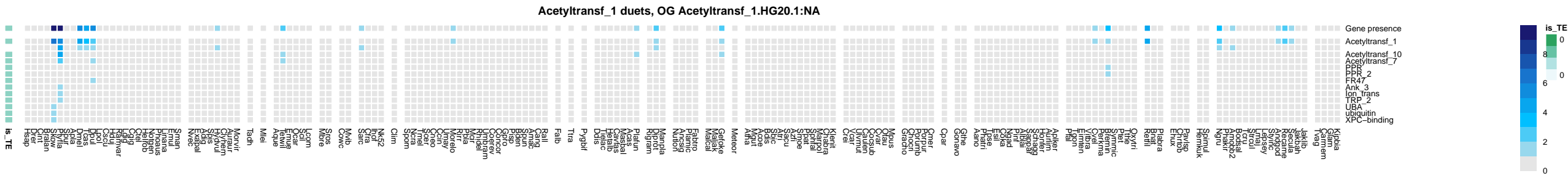




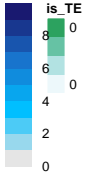








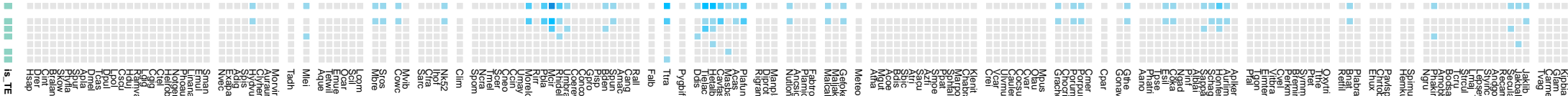


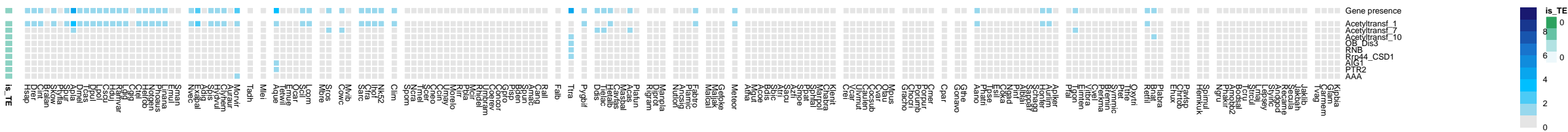


Gene presence

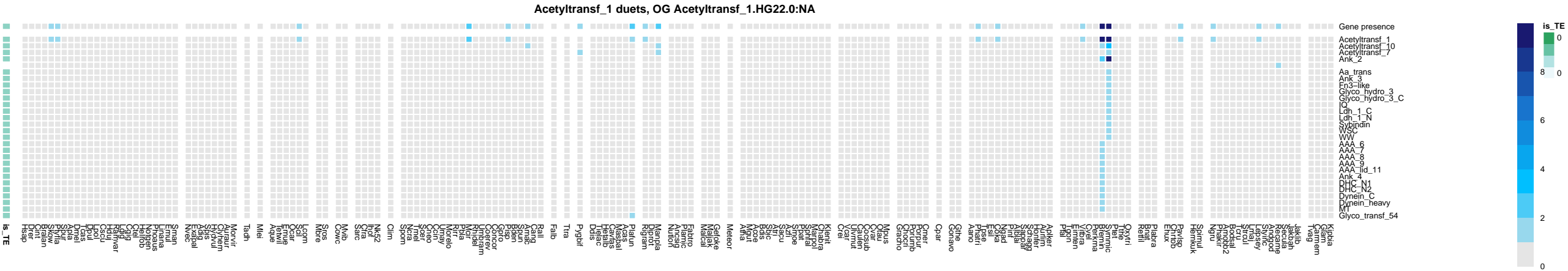
Acetyltransf\_1  
Acetyltransf\_10  
Acetyltransf\_7  
Acetyltransf\_9  
IPK

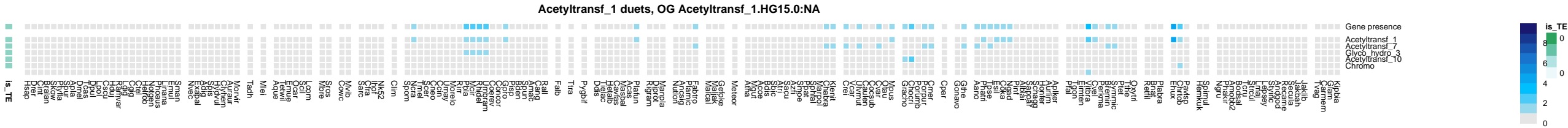
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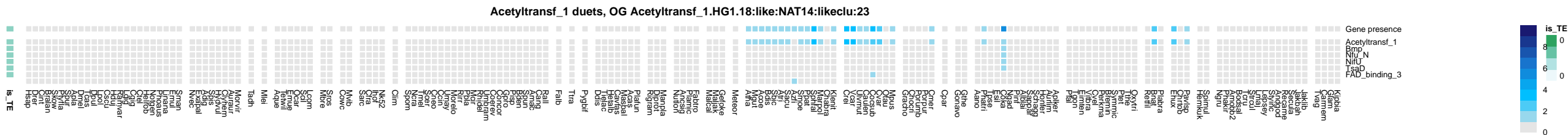


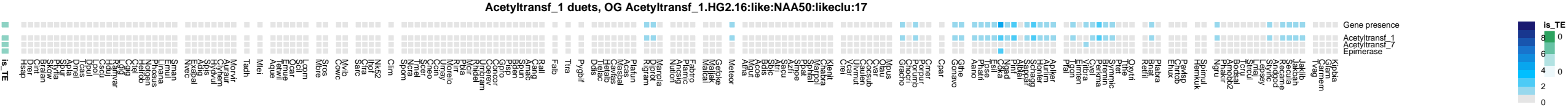


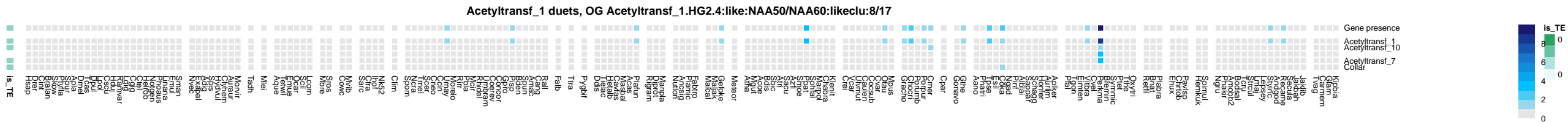
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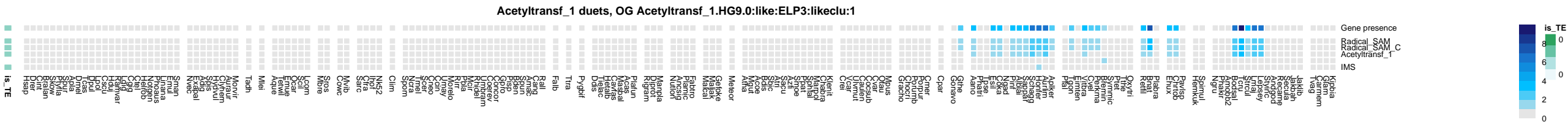




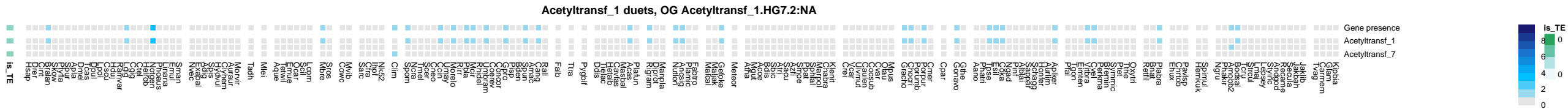


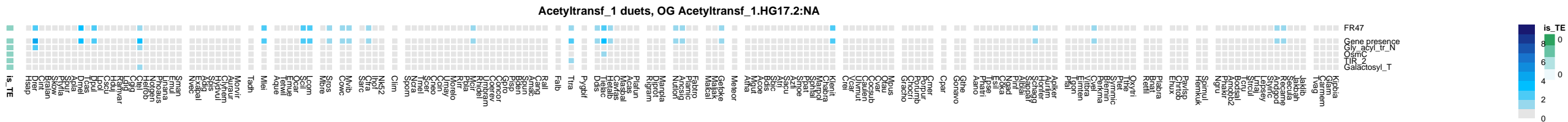


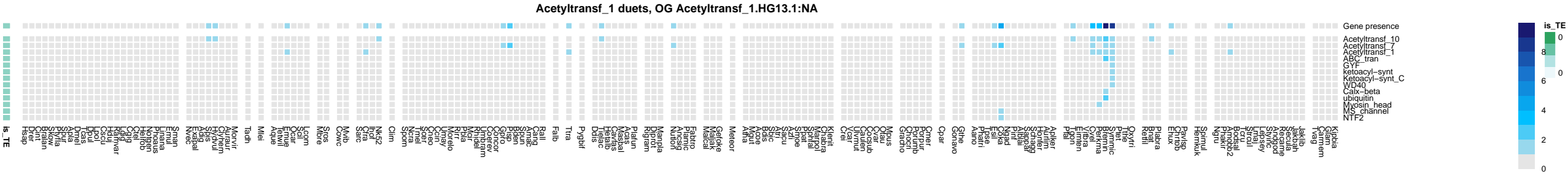


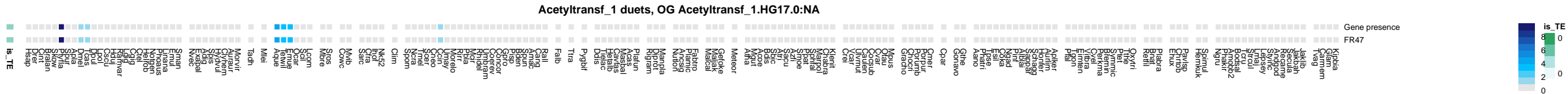


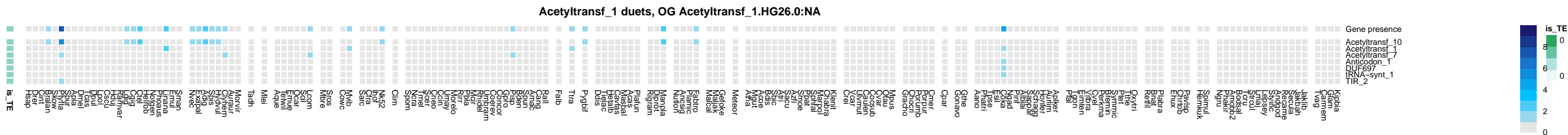


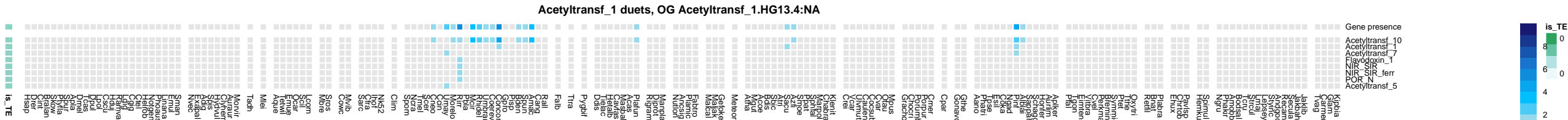




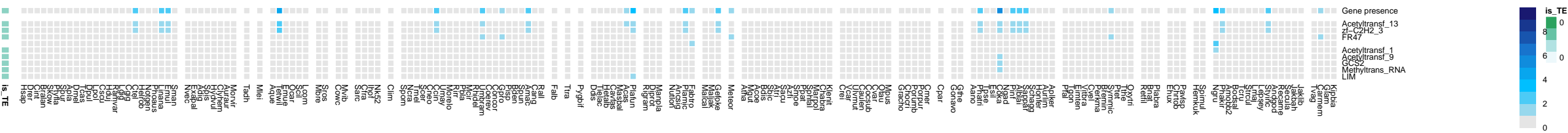


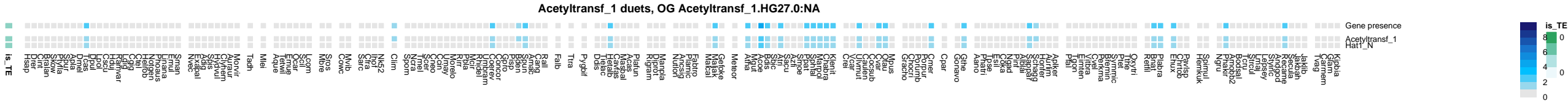




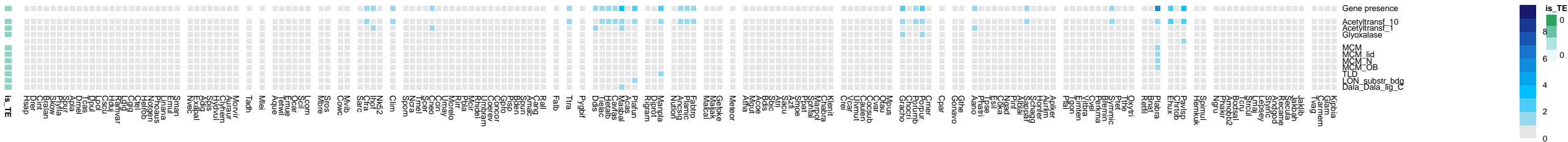


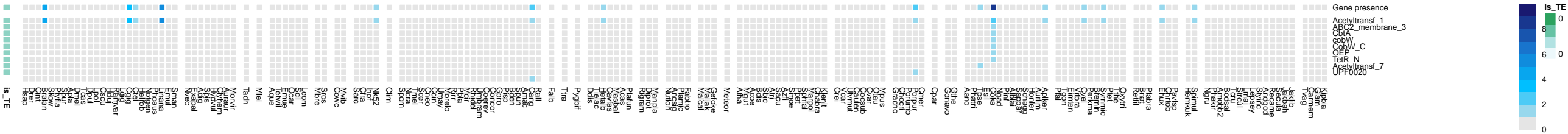
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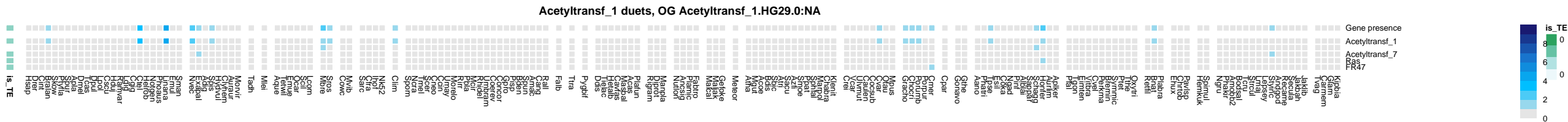


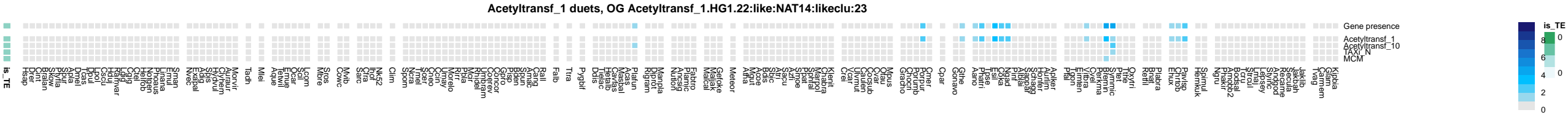


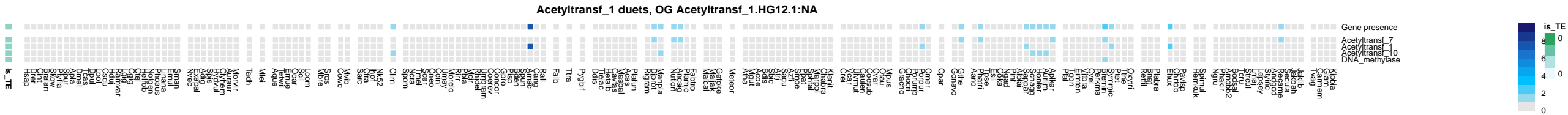


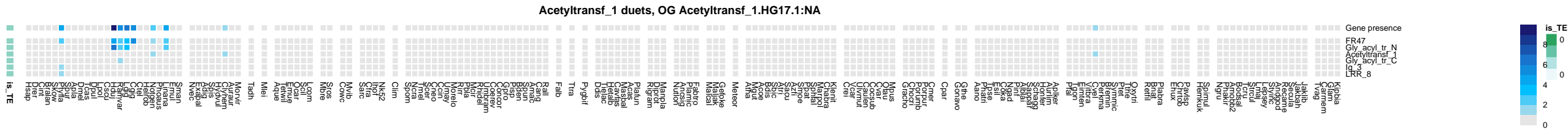


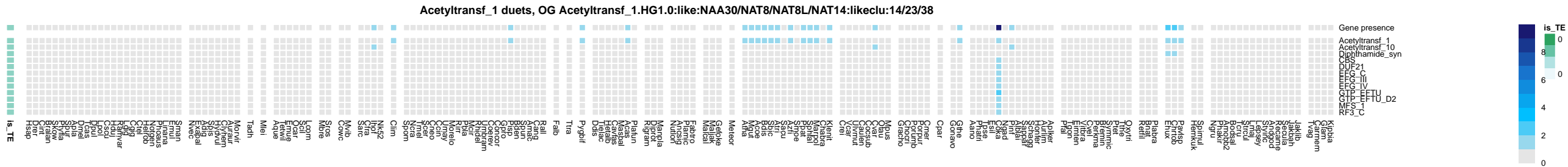


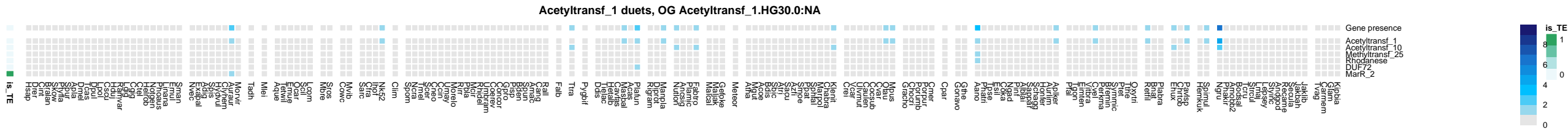




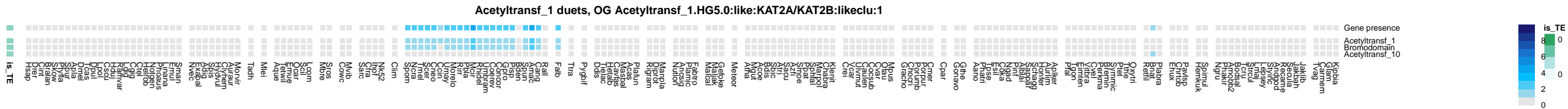


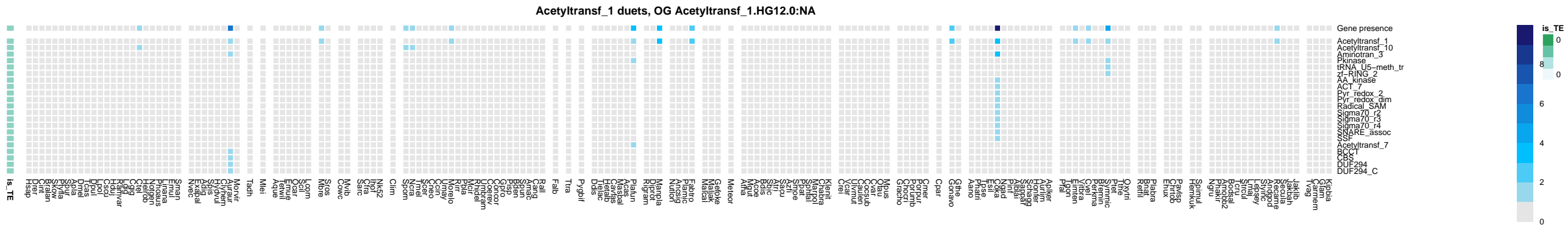


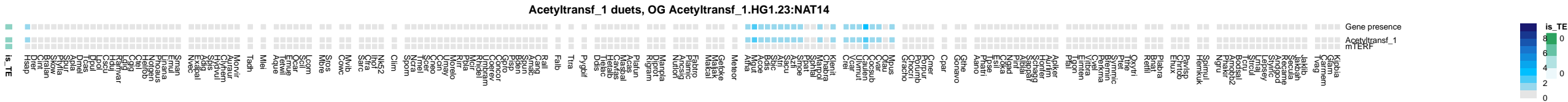


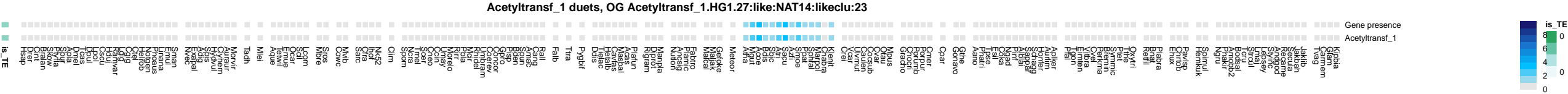


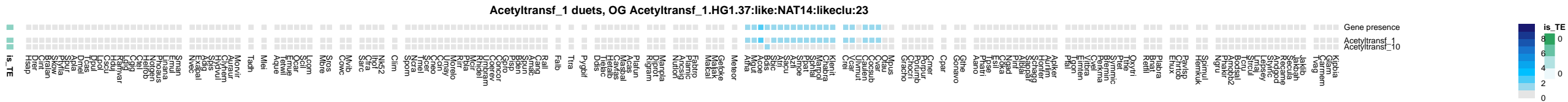


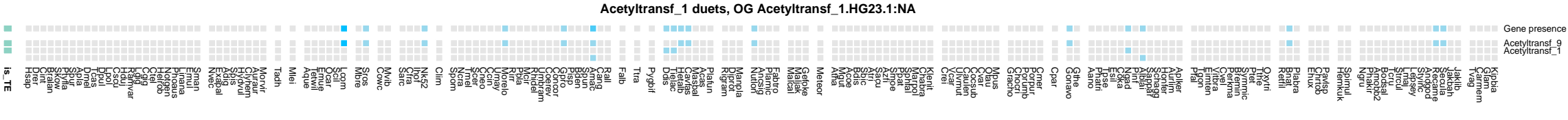
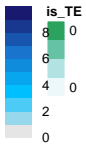


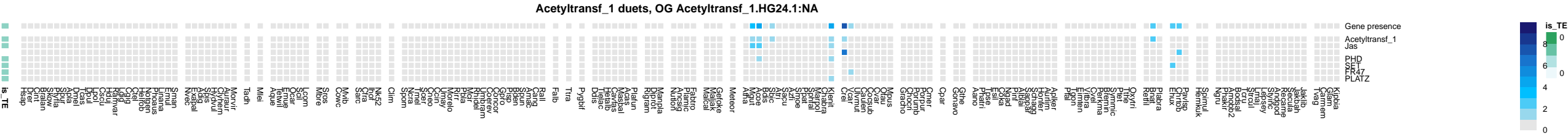




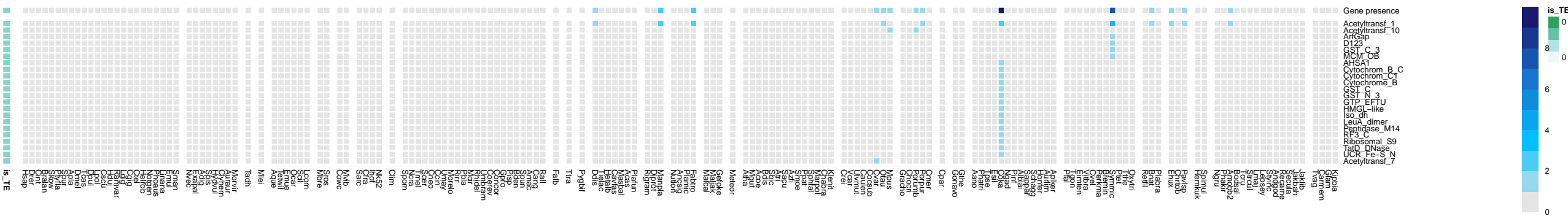




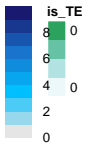




Acetyltransf\_1 duets, OG Acetyltransf\_1.HG1.2:like:NAA30/NAT8/NAT8L/NAT14:likeclu:14/23/38



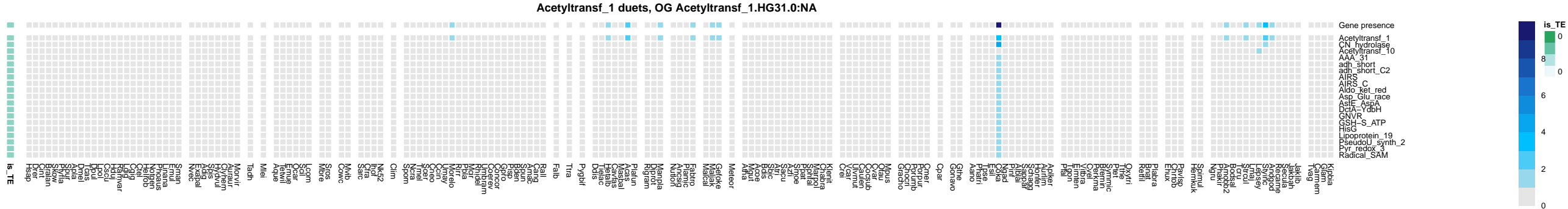


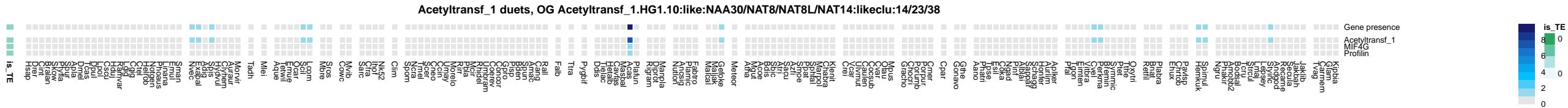


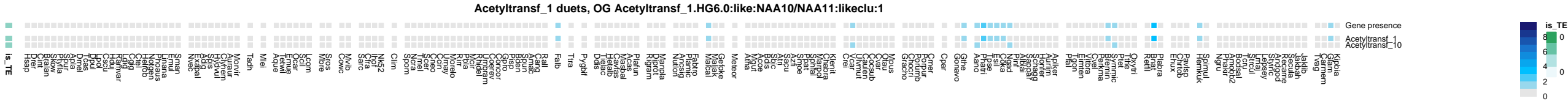
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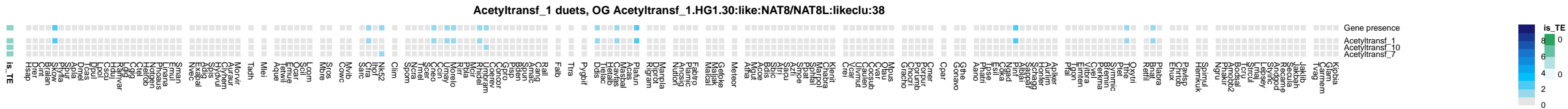


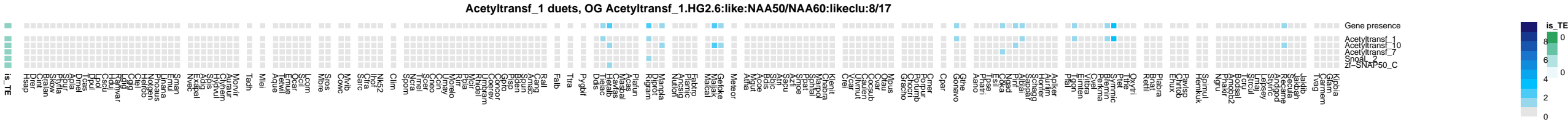


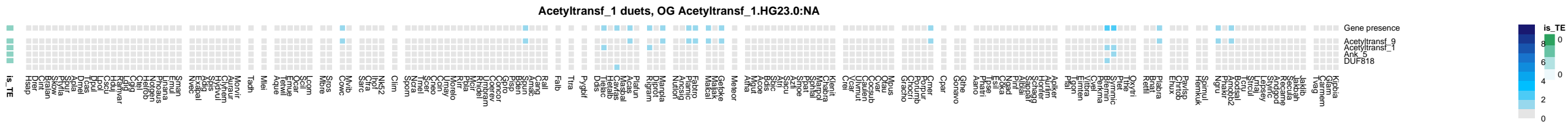




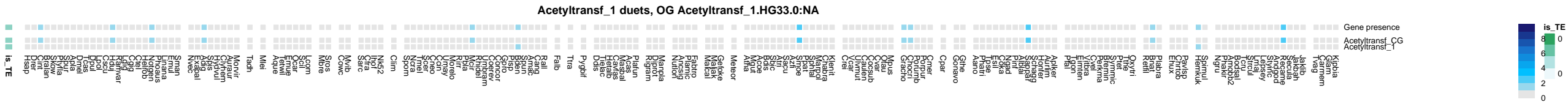


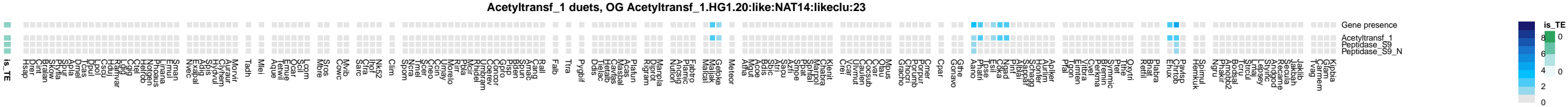




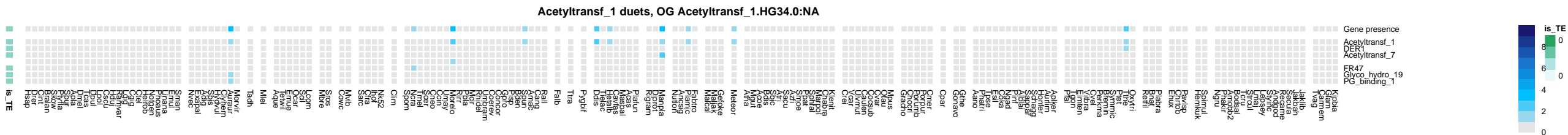


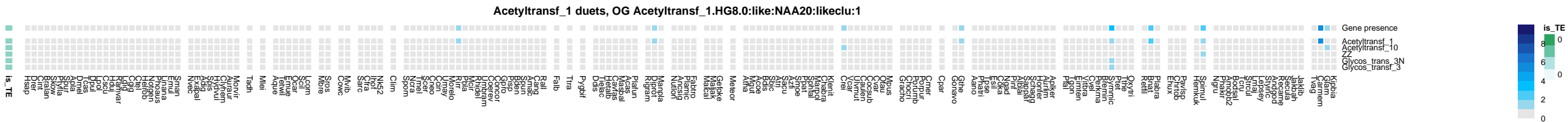


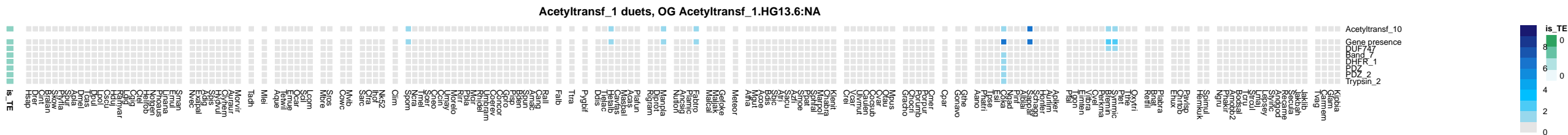


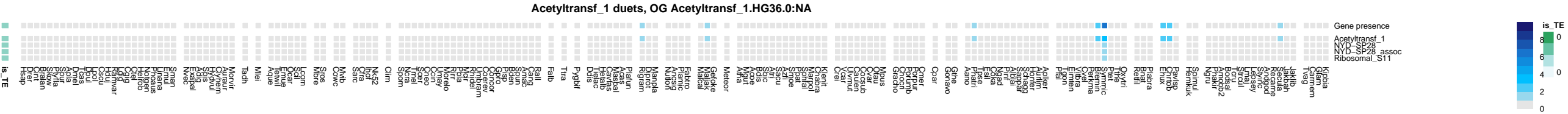


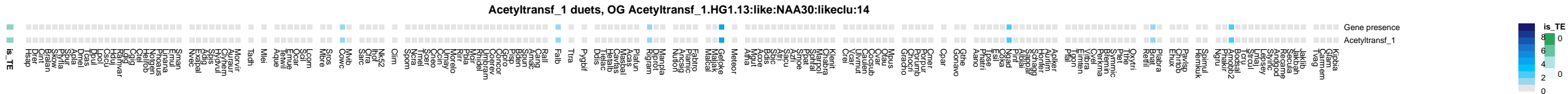




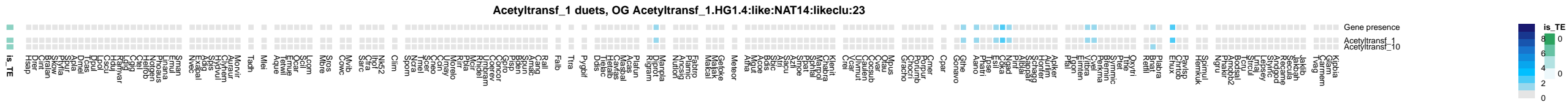


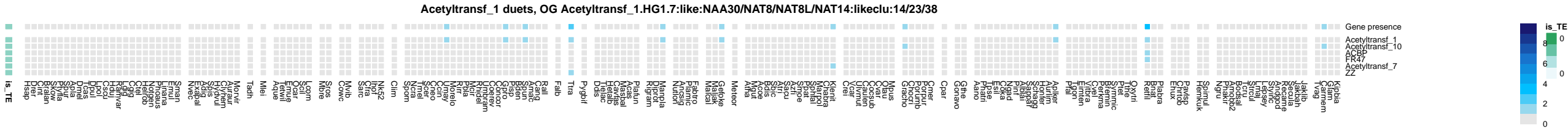


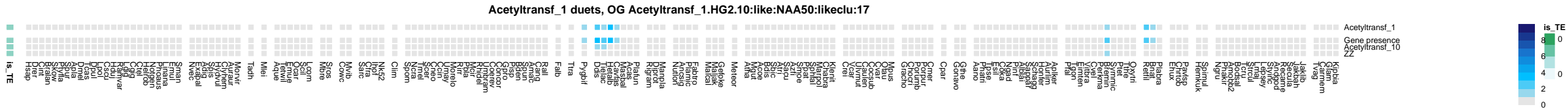


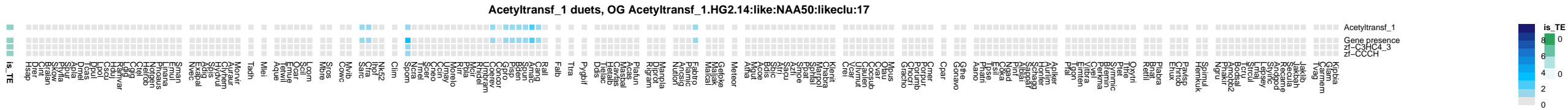


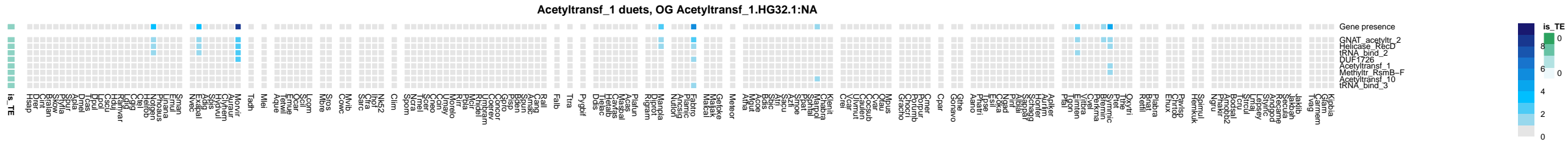


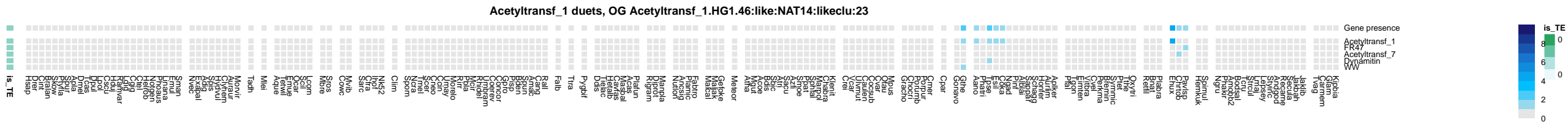


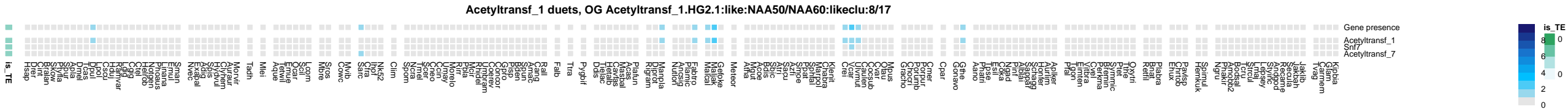


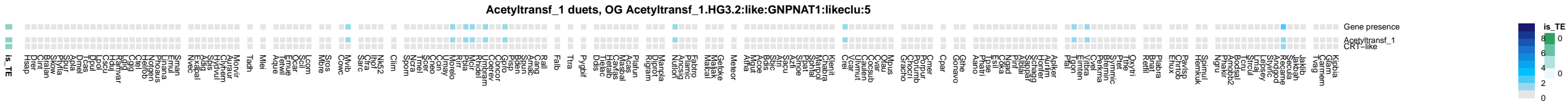




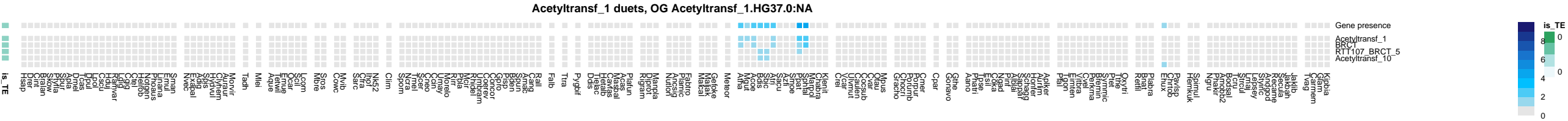


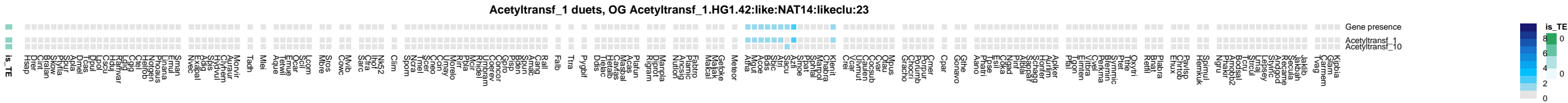


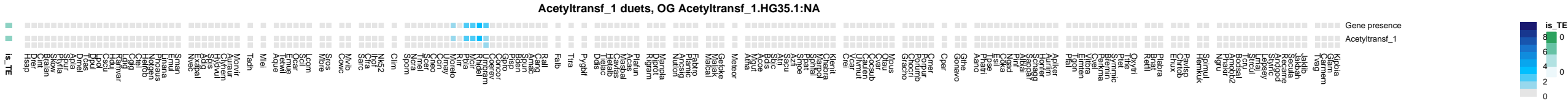


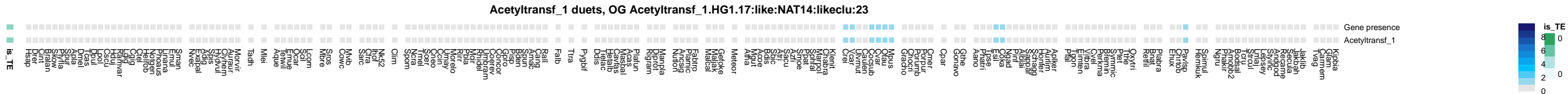


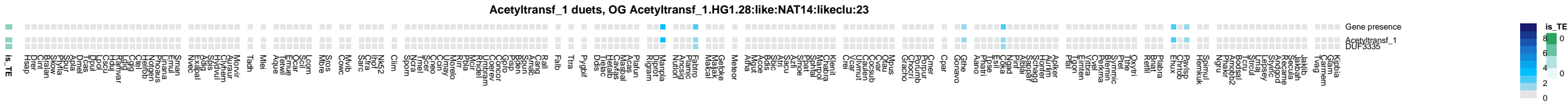


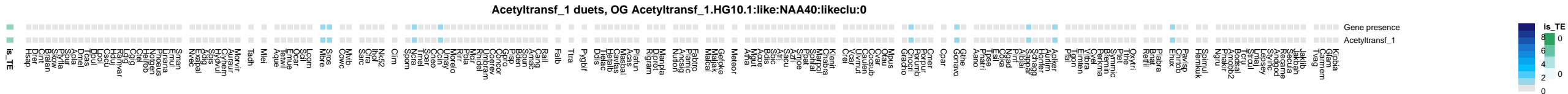


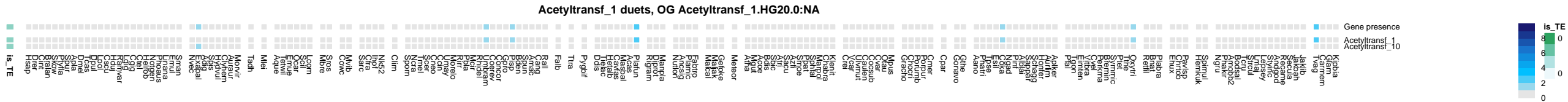


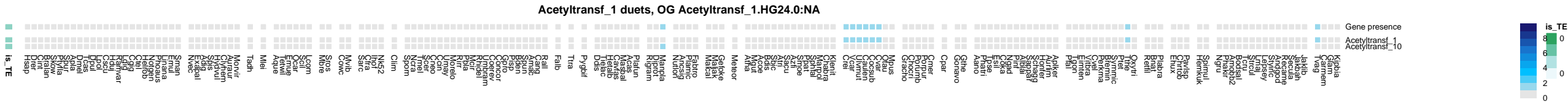




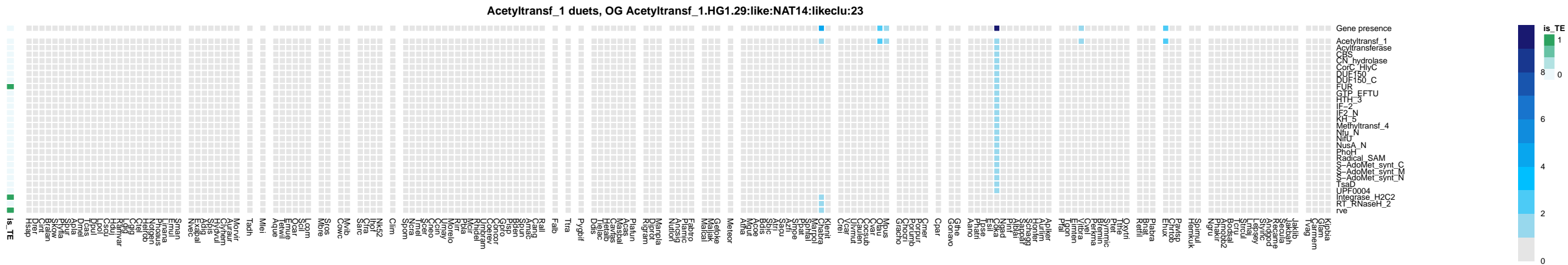


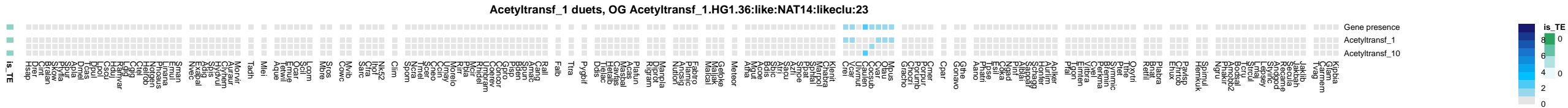


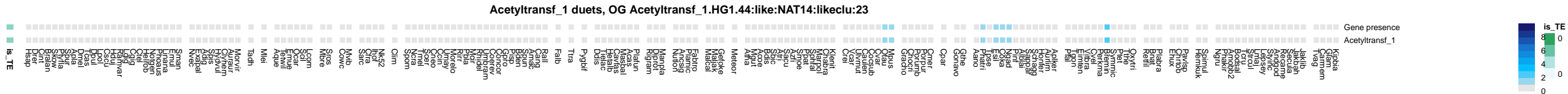


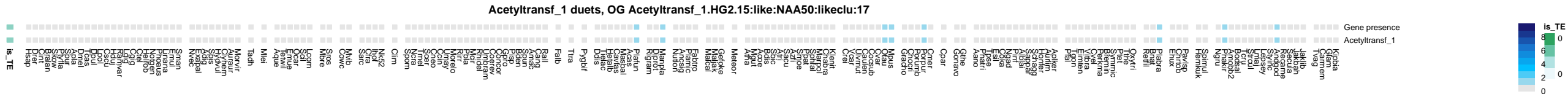


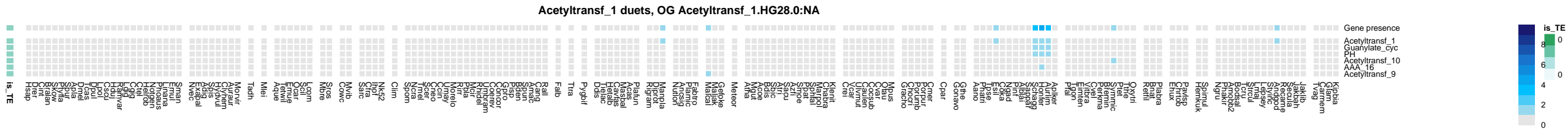


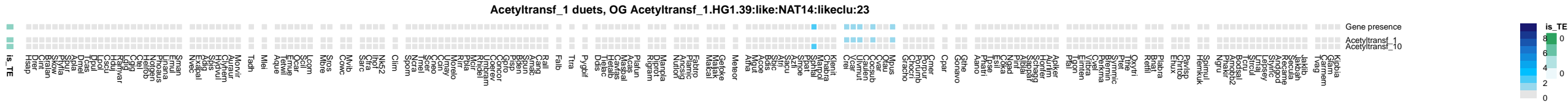


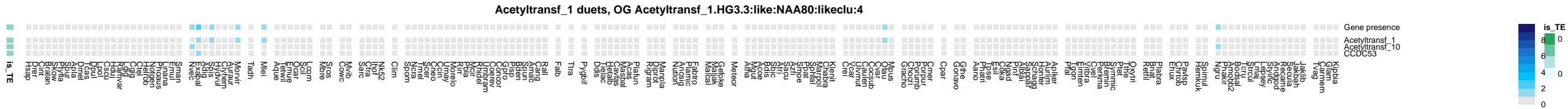


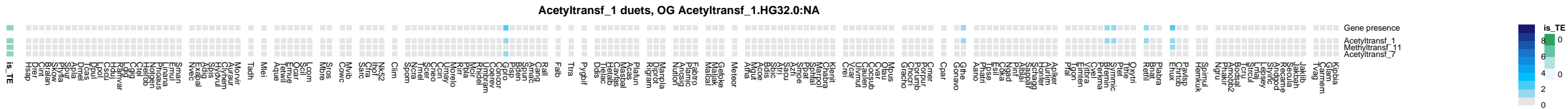




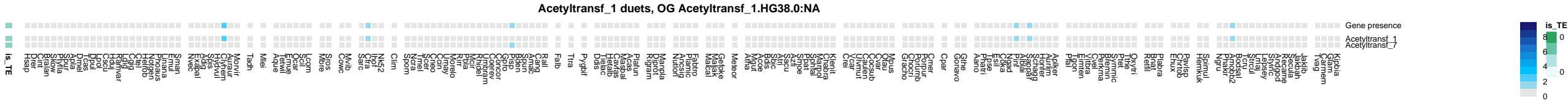


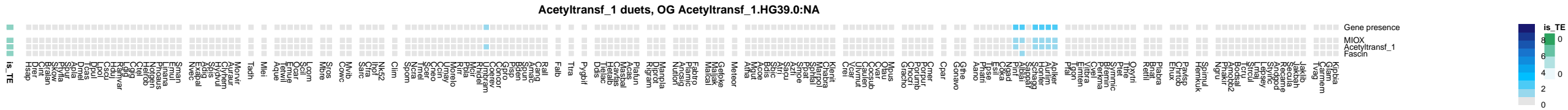


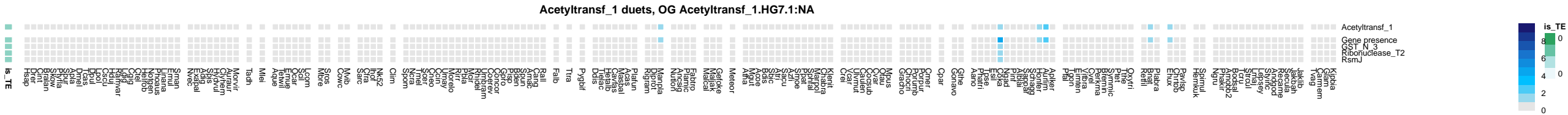


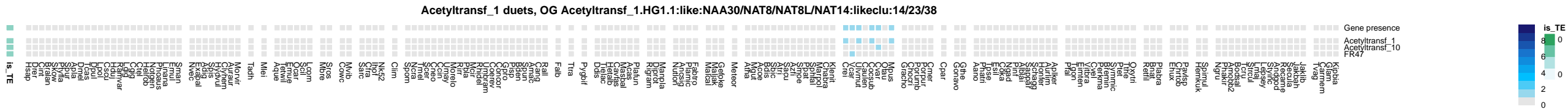




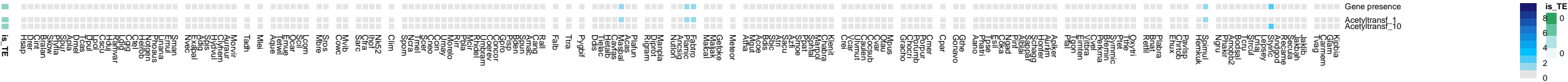


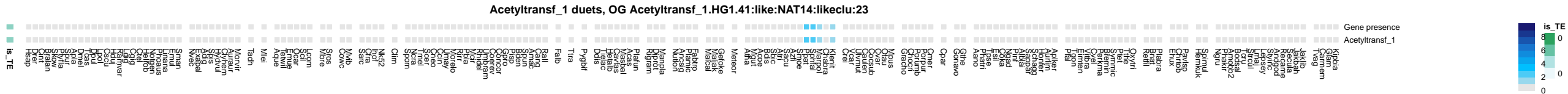


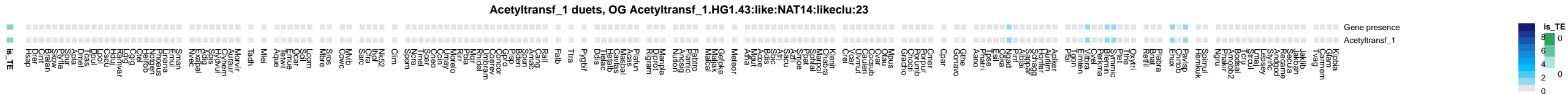


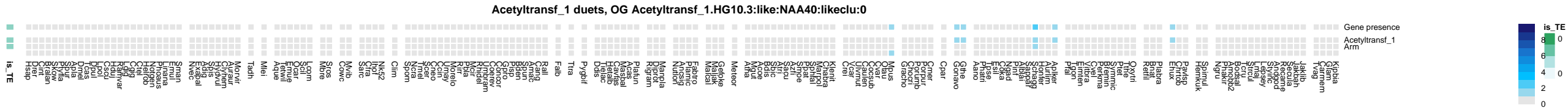


Acetyltransf\_1 duets, OG Acetyltransf\_1.HG1.11:like:NAA30/NAT8L/NAT14:liikeclu:14/23/38

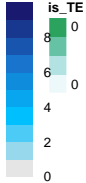












Acetyltransf\_1 duets, OG Acetyltransf\_1.HG13.2:NA



