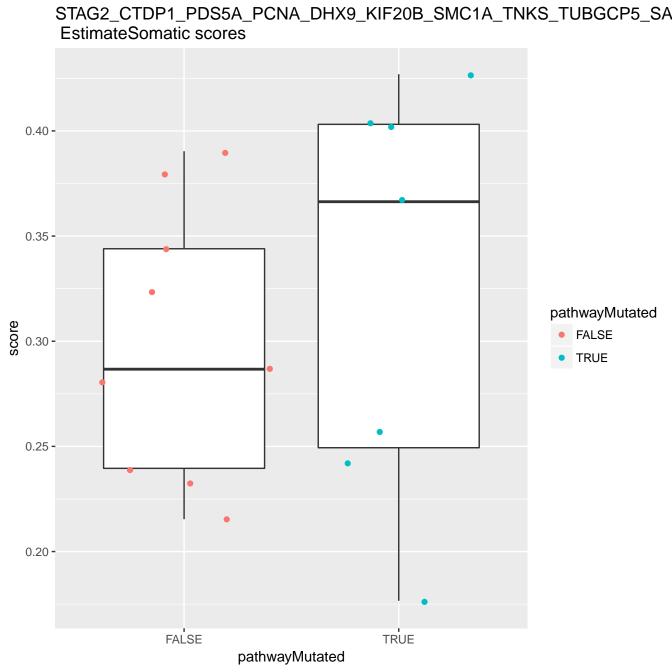
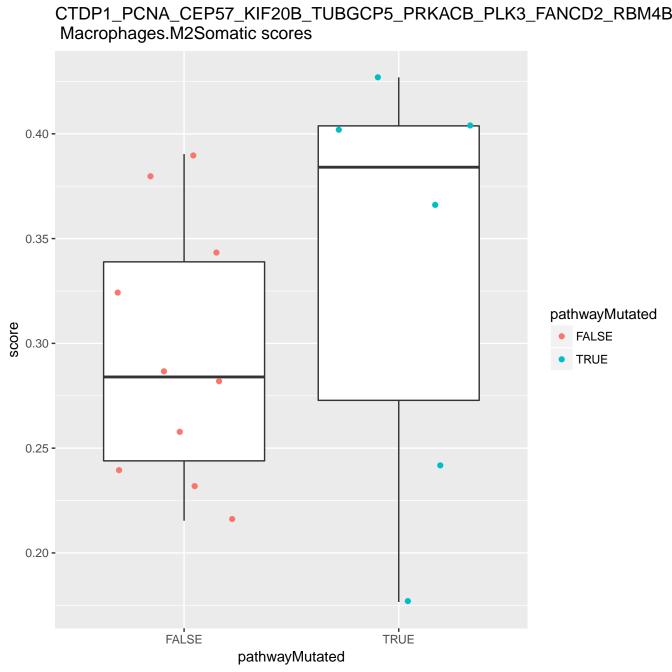
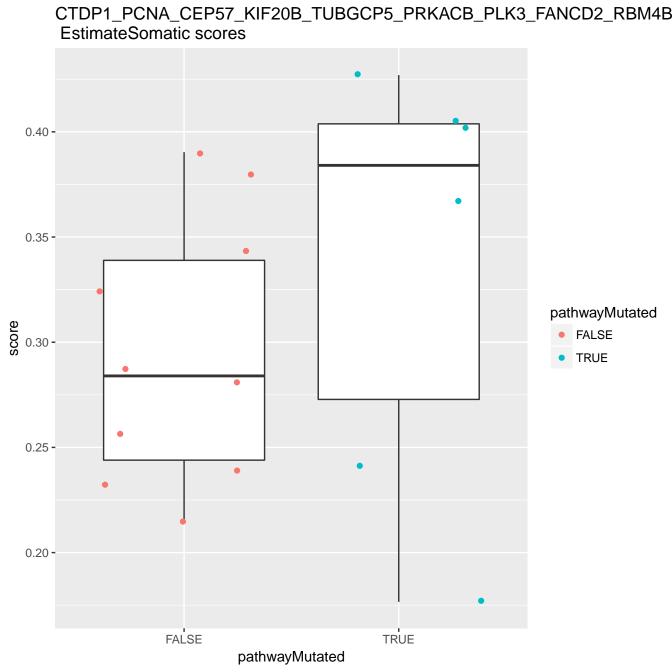


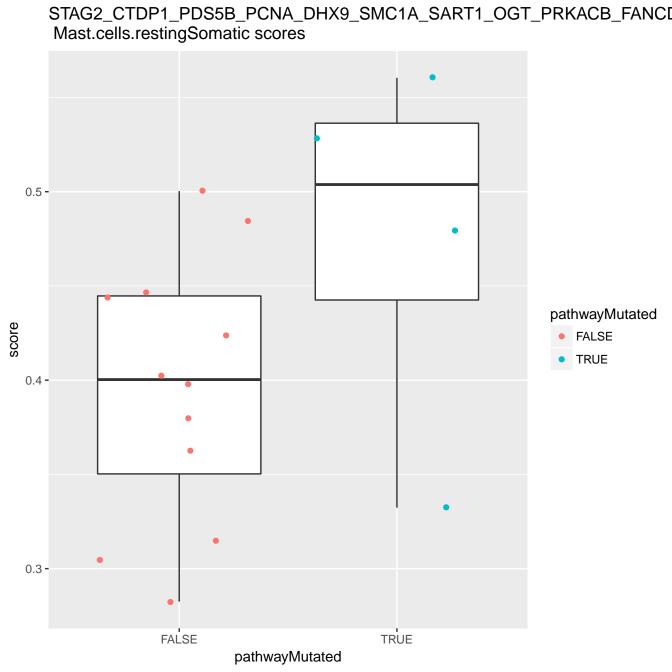
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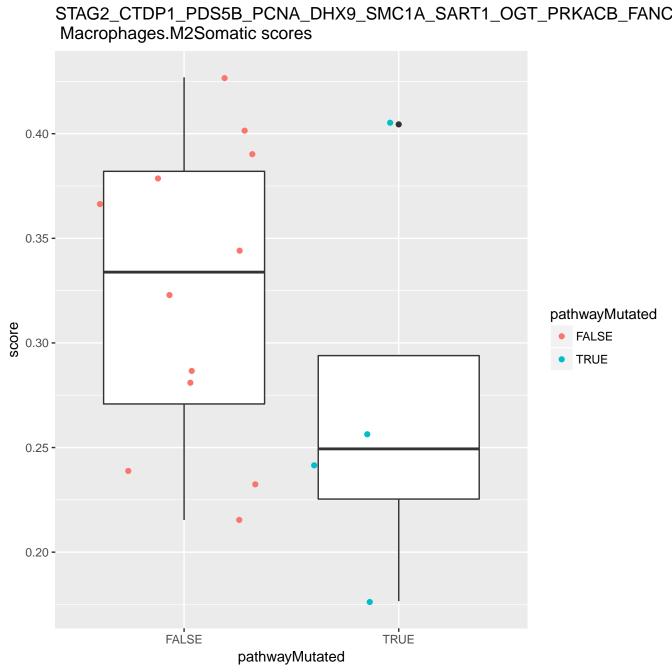


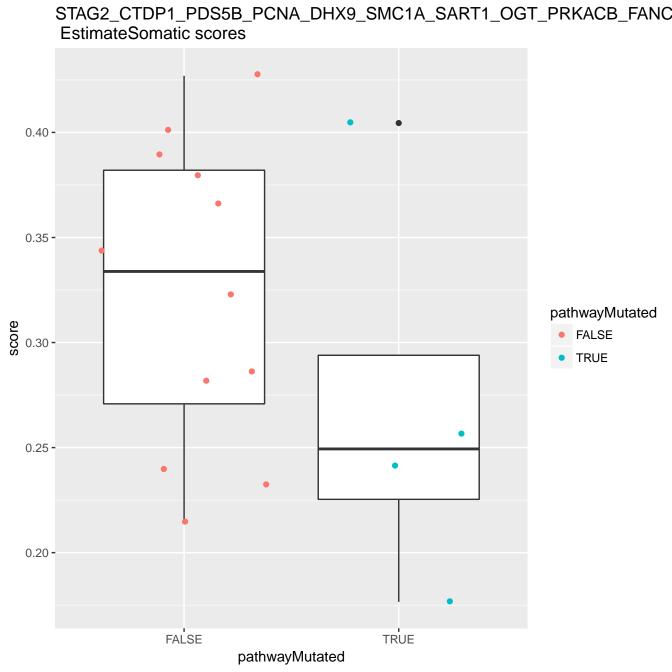
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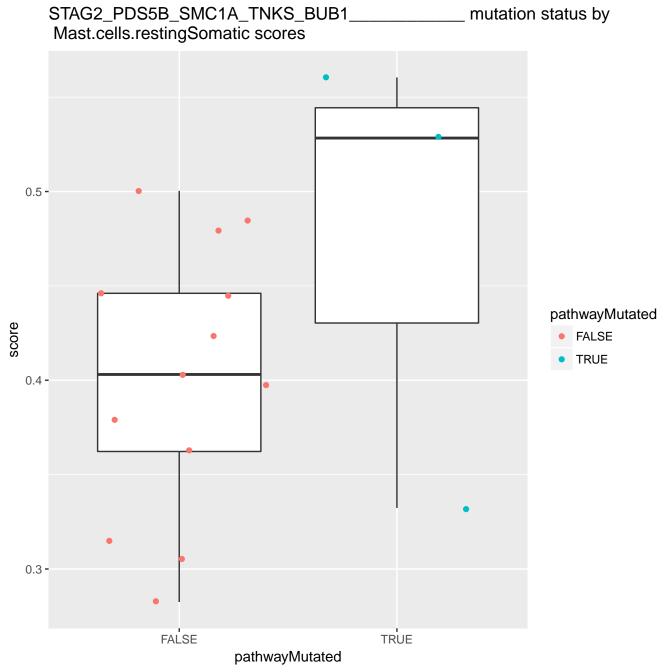


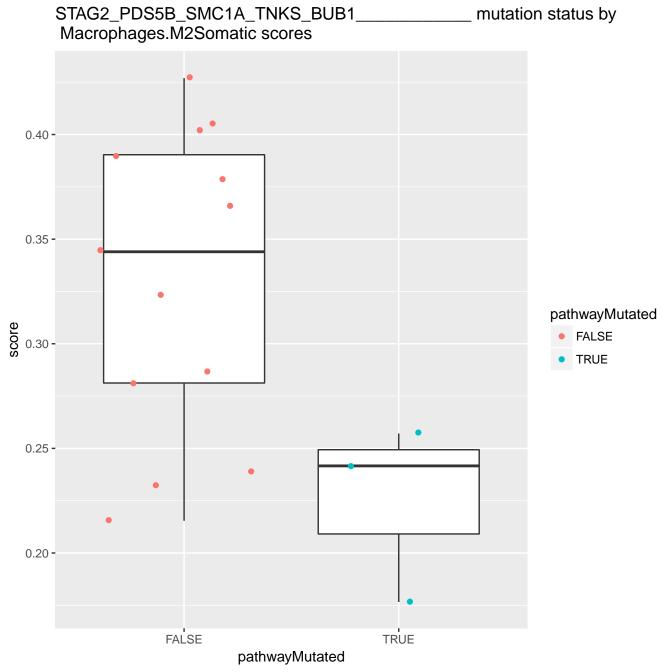


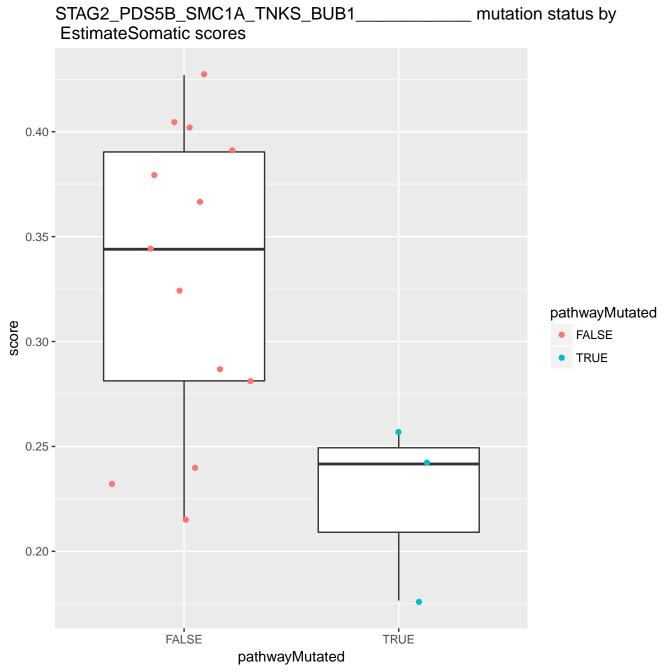


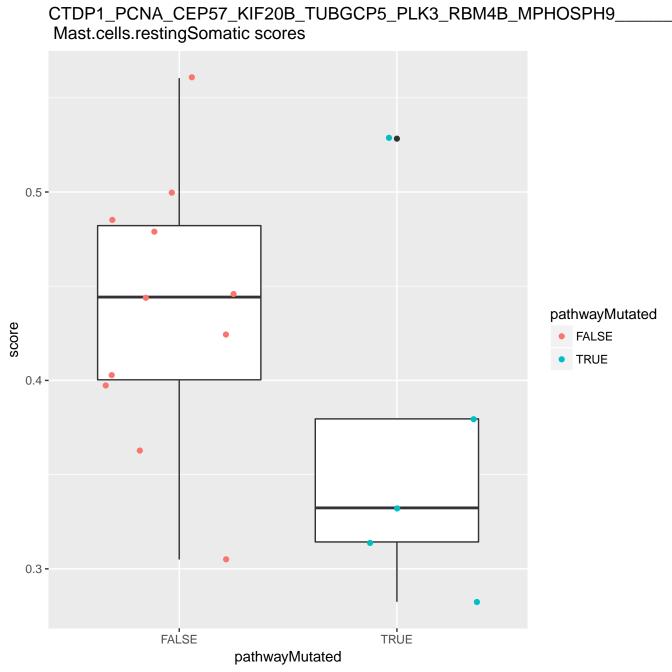


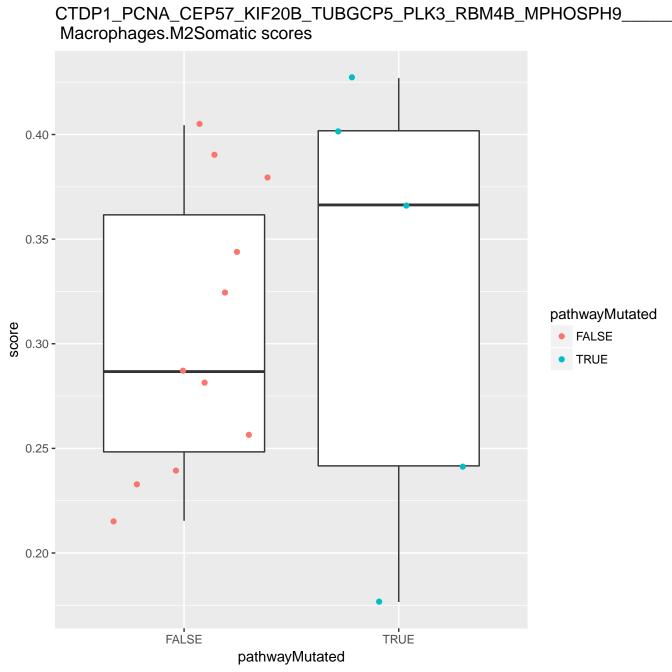


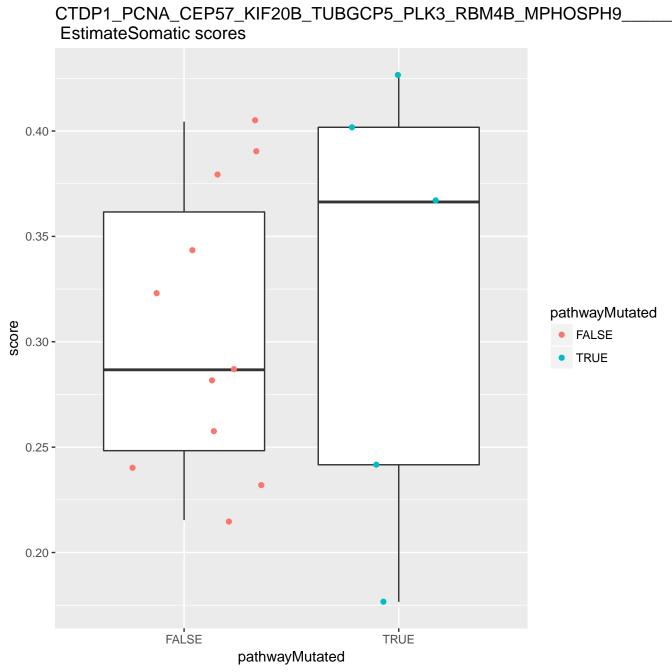


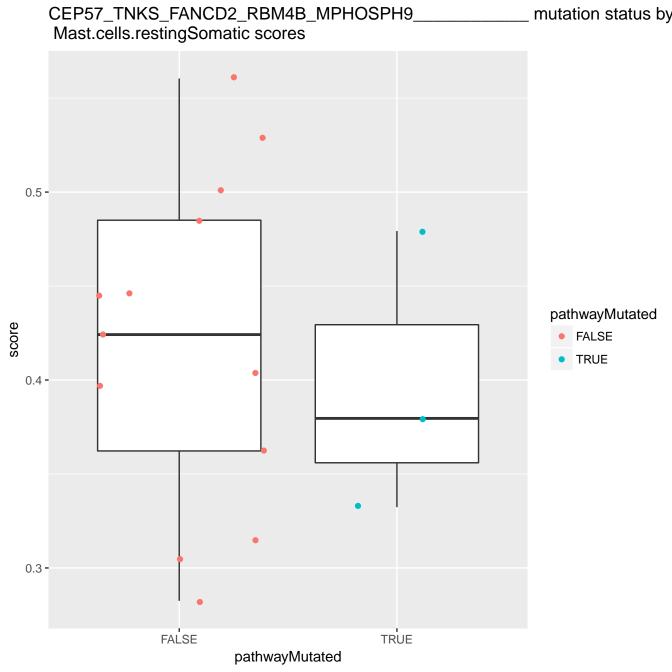


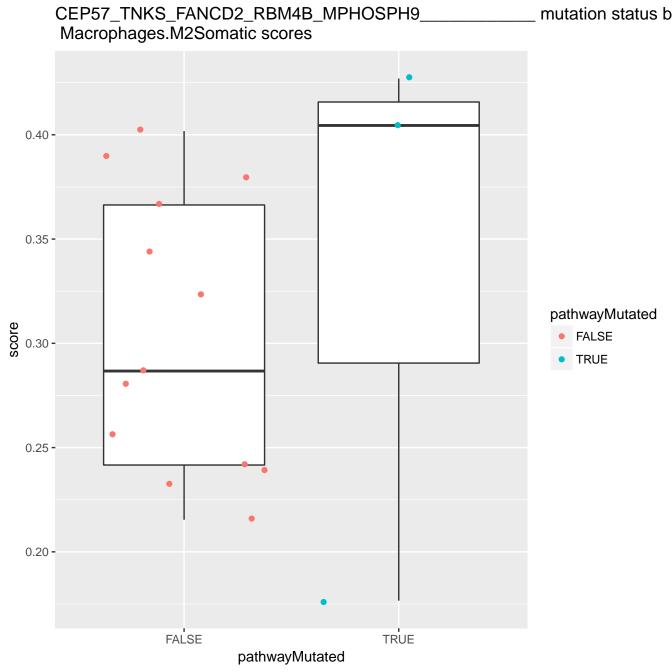


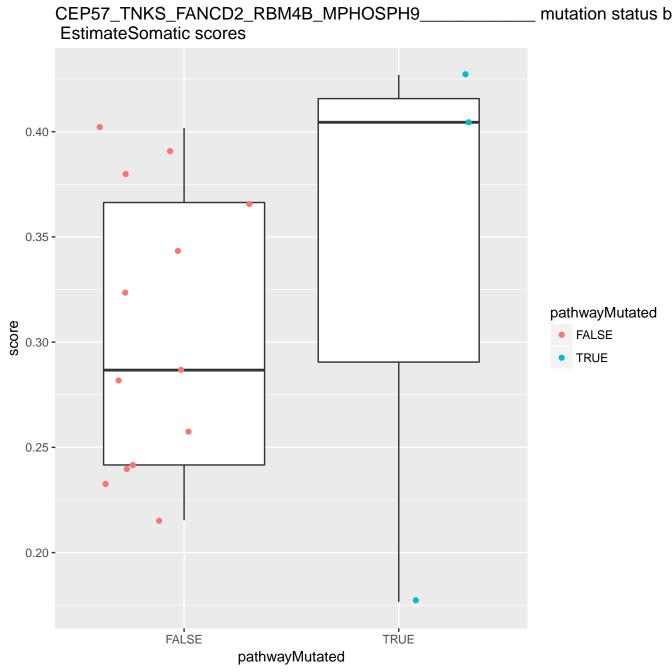


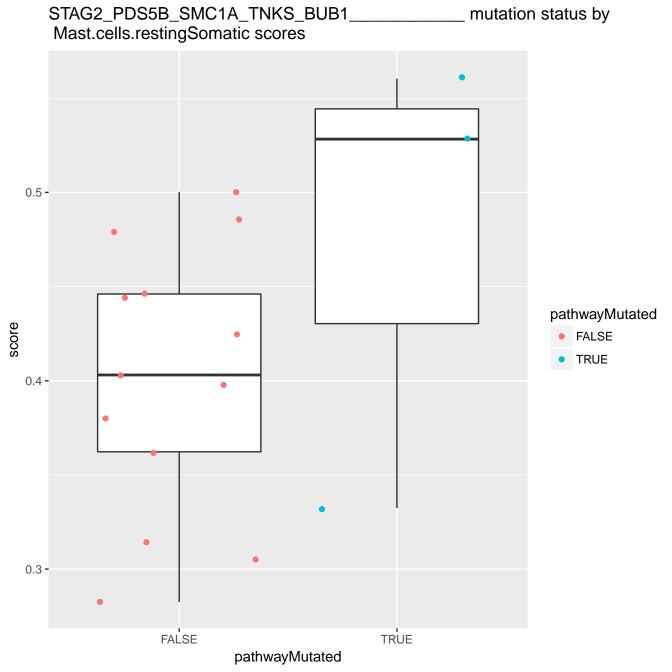


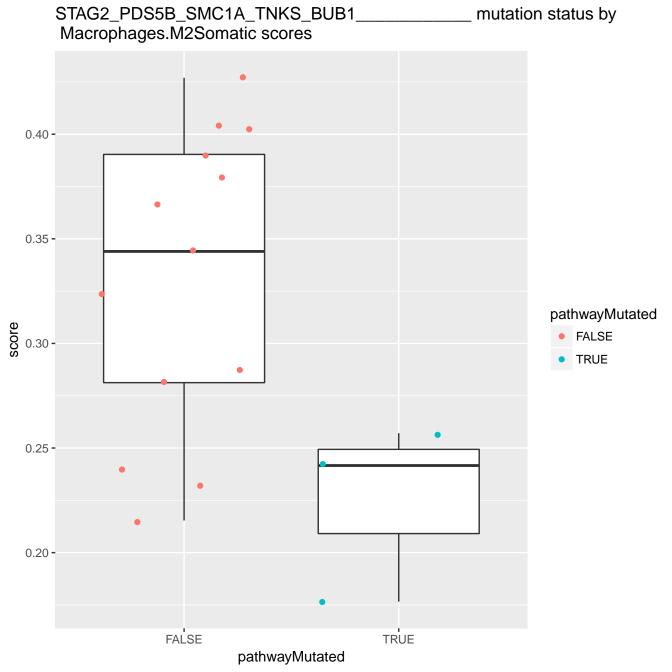


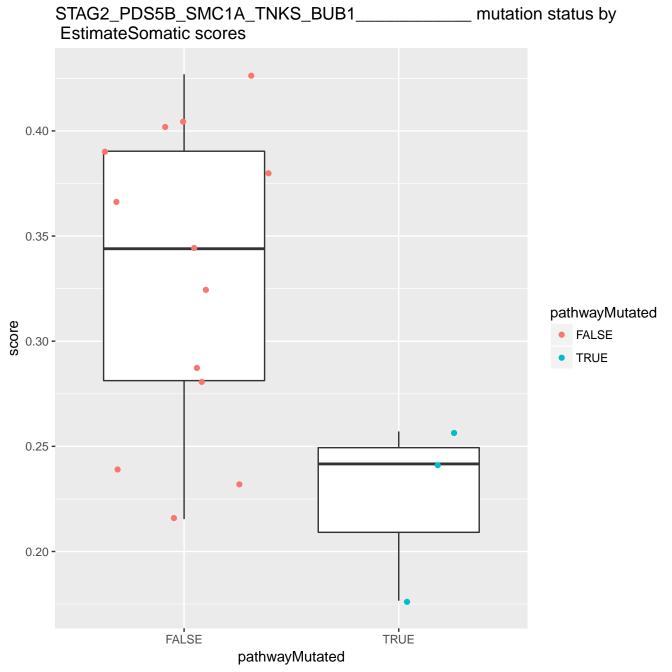


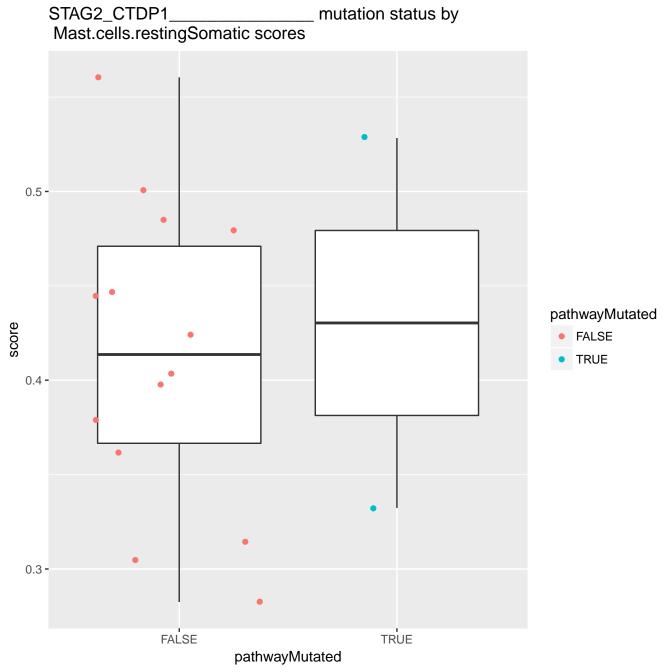


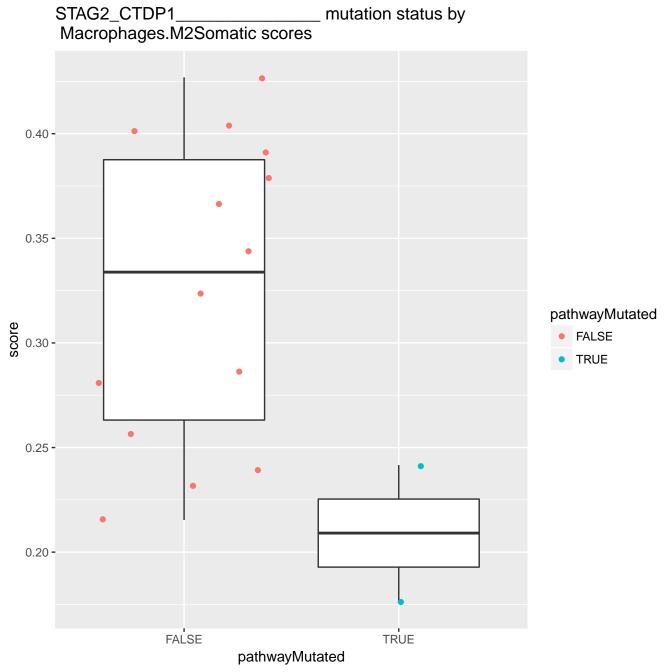


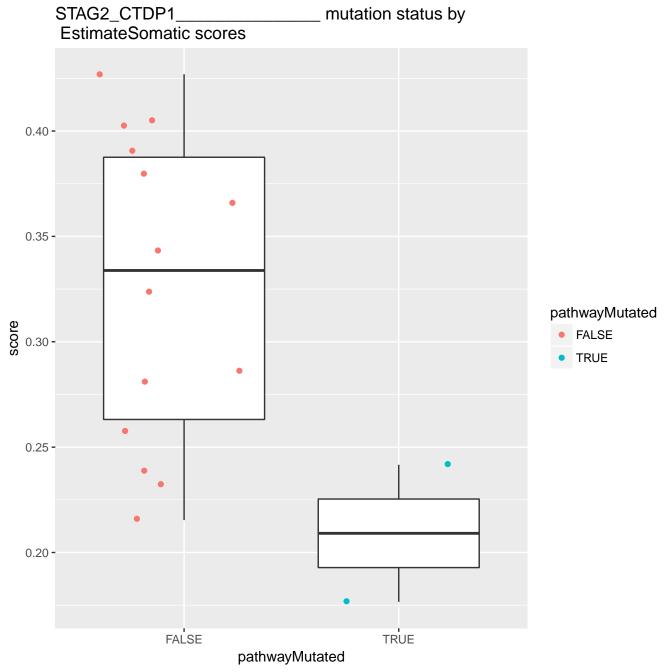


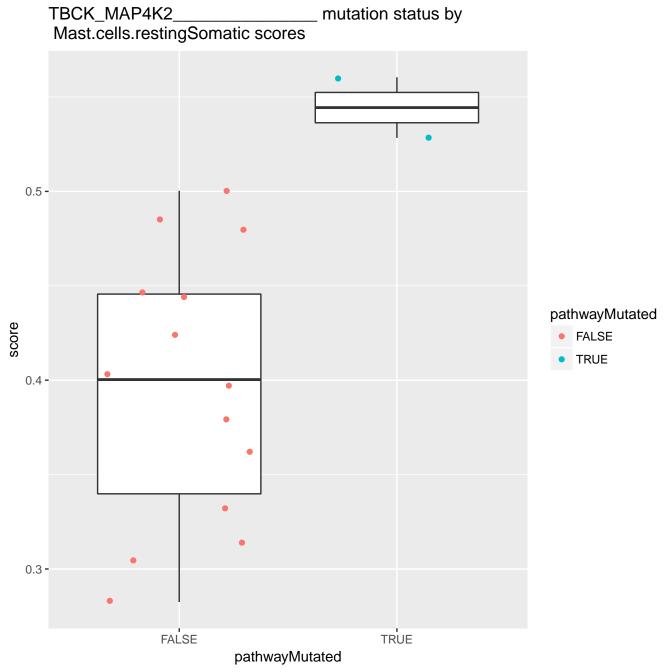


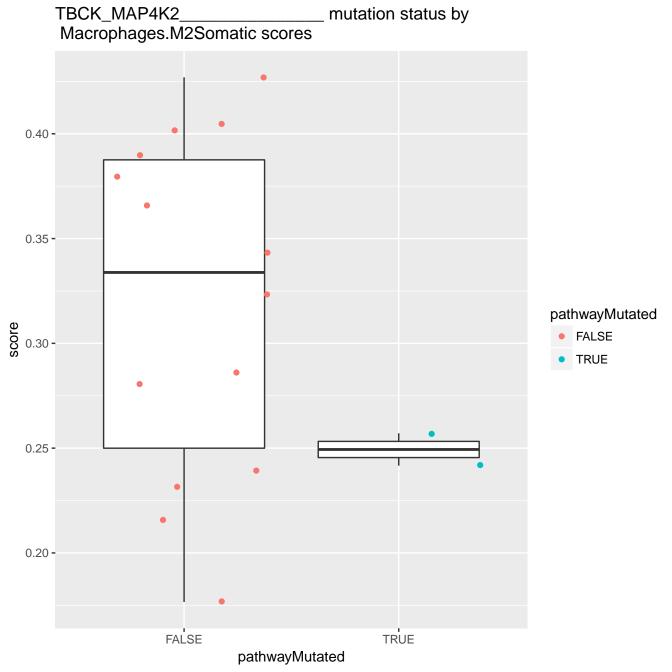


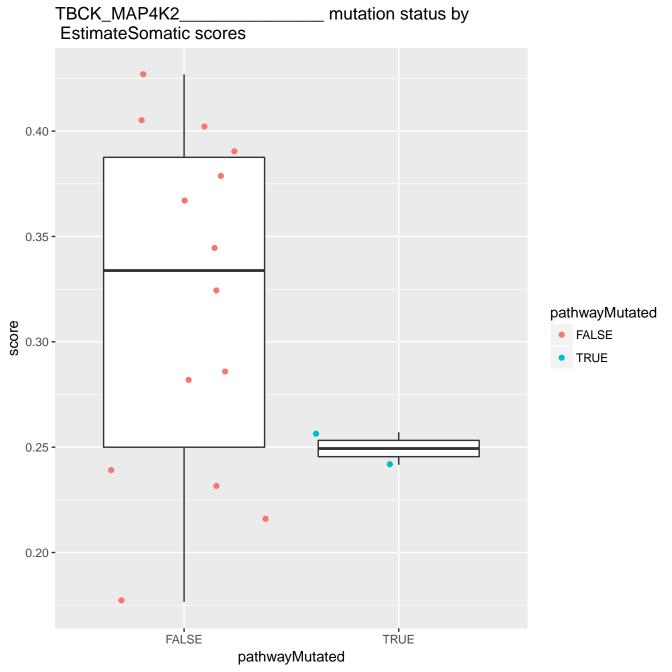


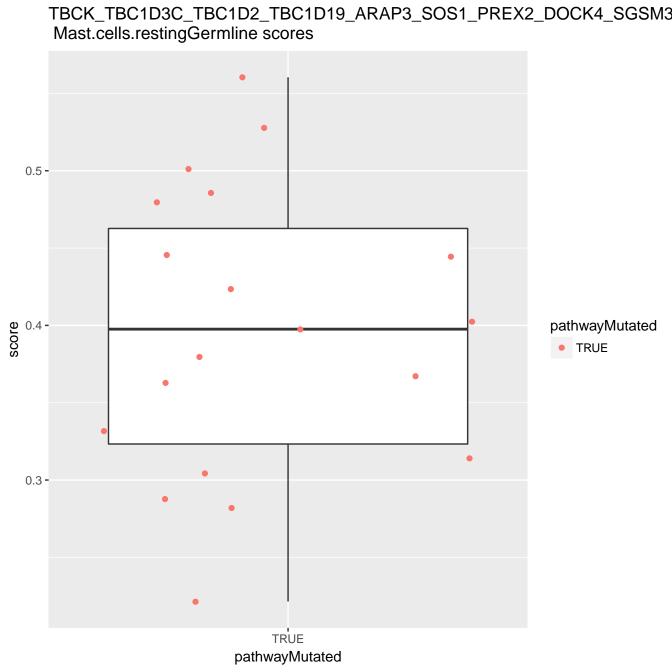




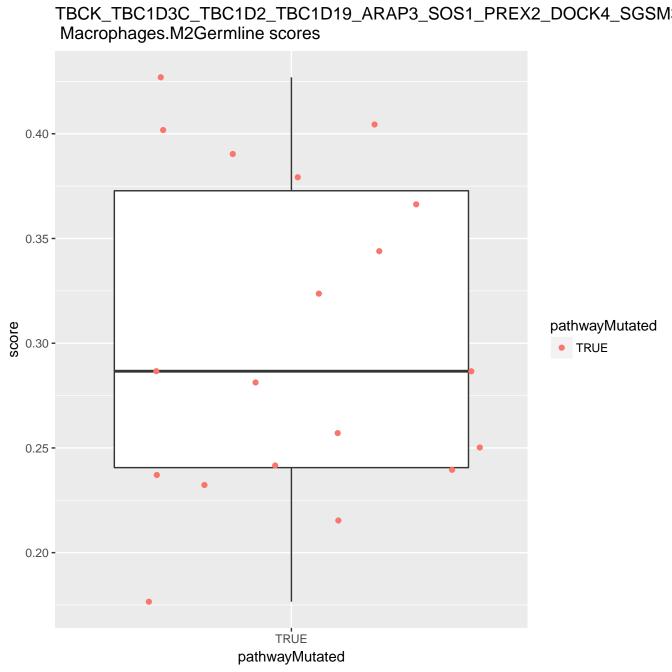




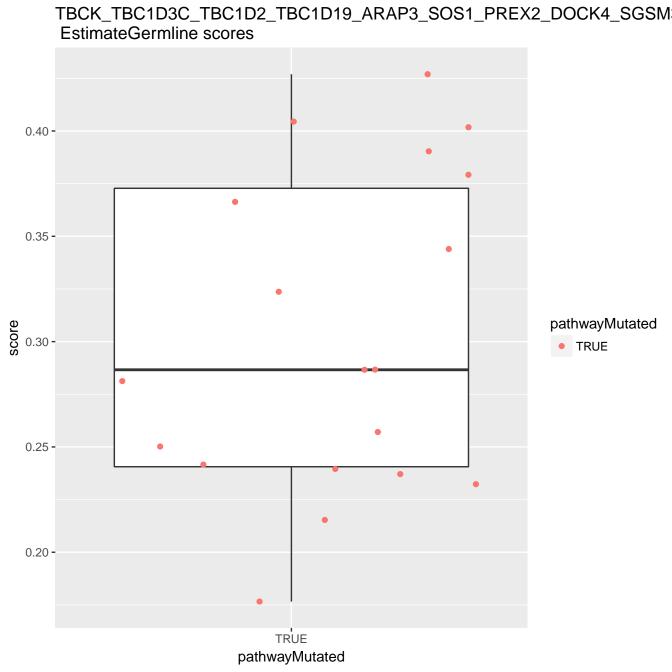


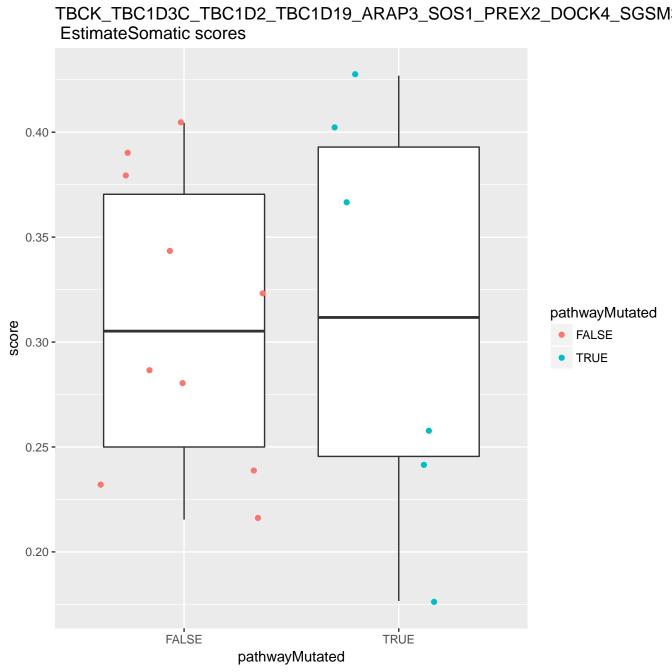


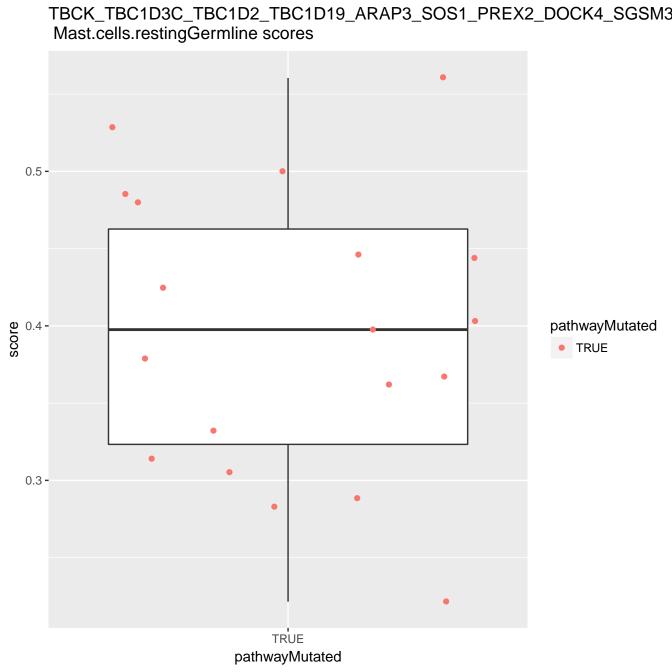
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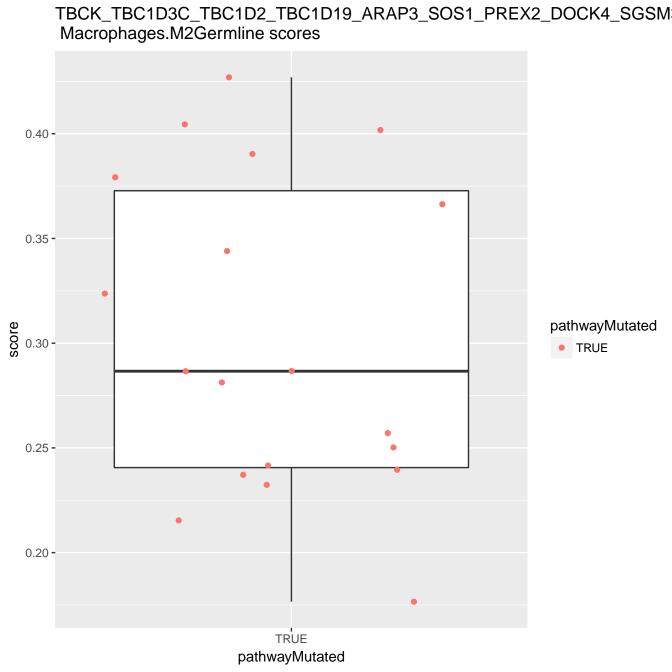
TBCK\_TBC1D3C\_TBC1D2\_TBC1D19\_ARAP3\_SOS1\_PREX2\_DOCK4\_SGSM Macrophages.M2Somatic scores 0.40 -0.35 pathwayMutated 800 -0.30 -FALSE TRUE 0.25 -0.20 -FALSE TRUE pathwayMutated



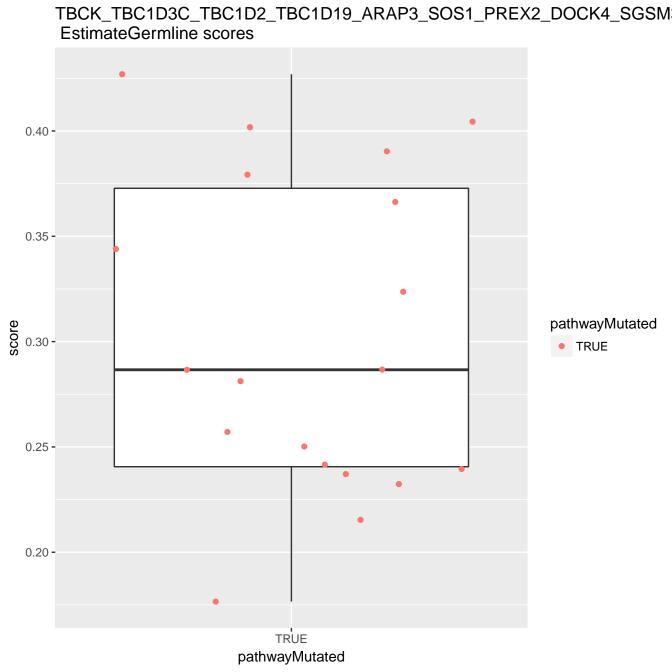


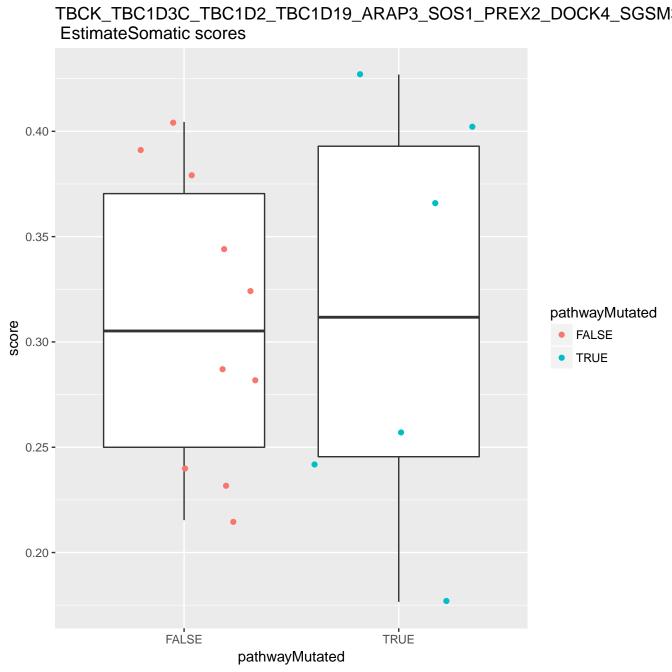


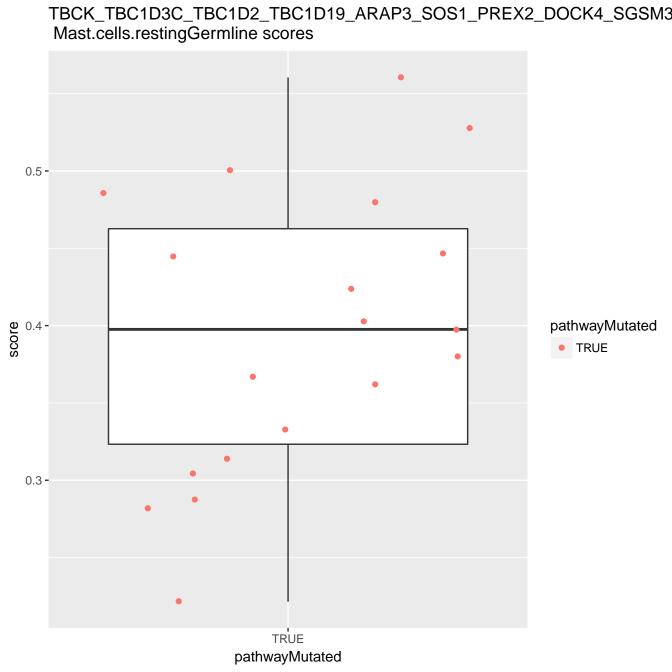
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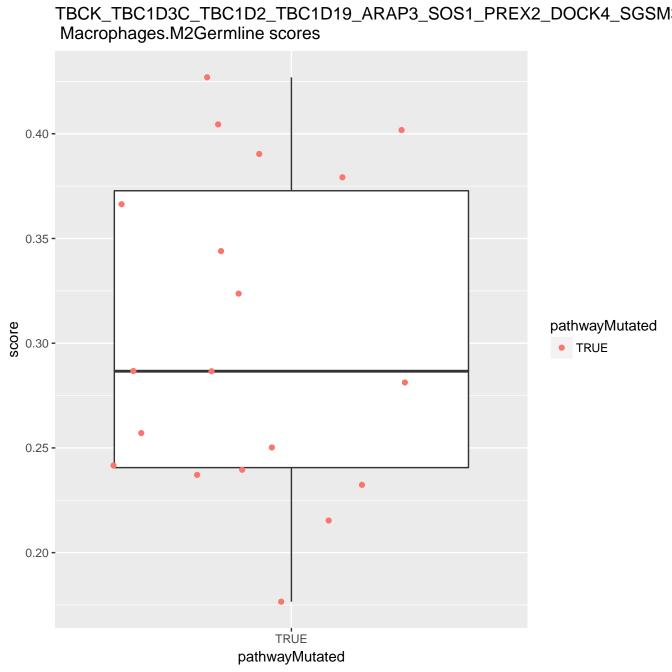
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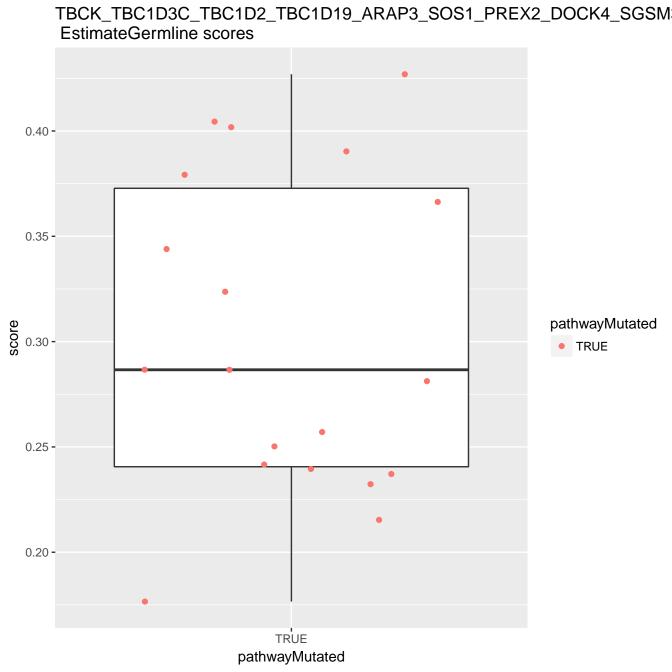


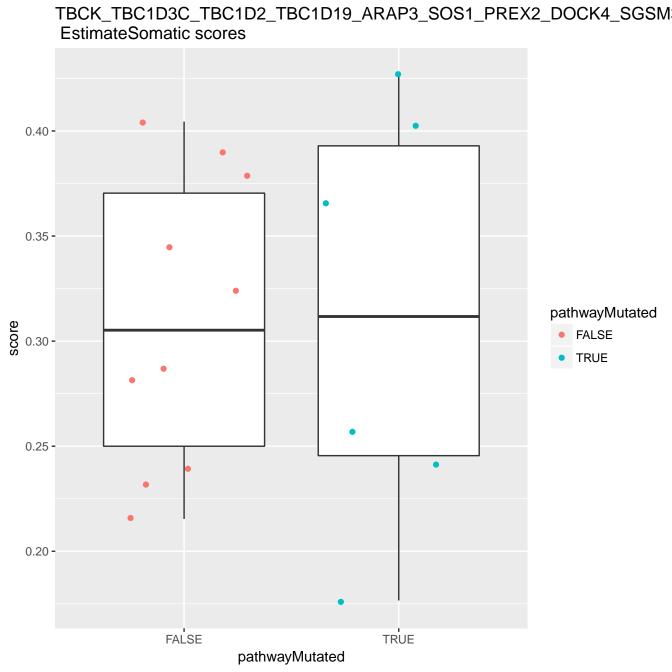


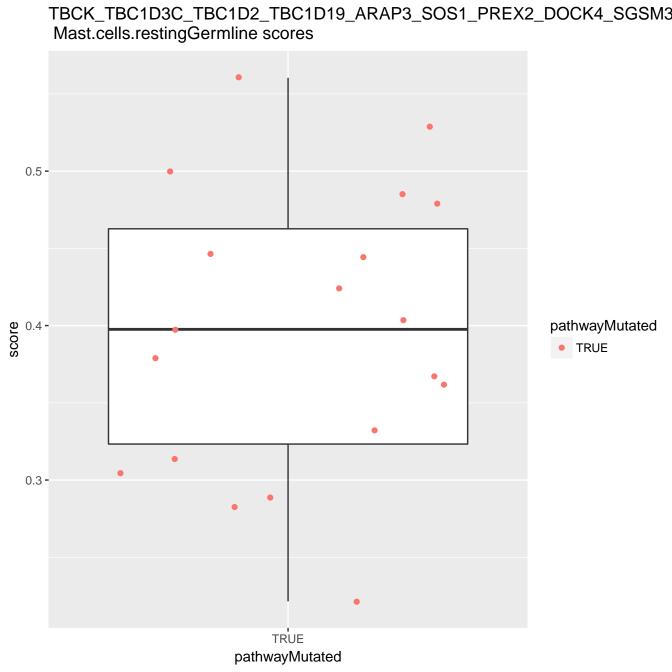
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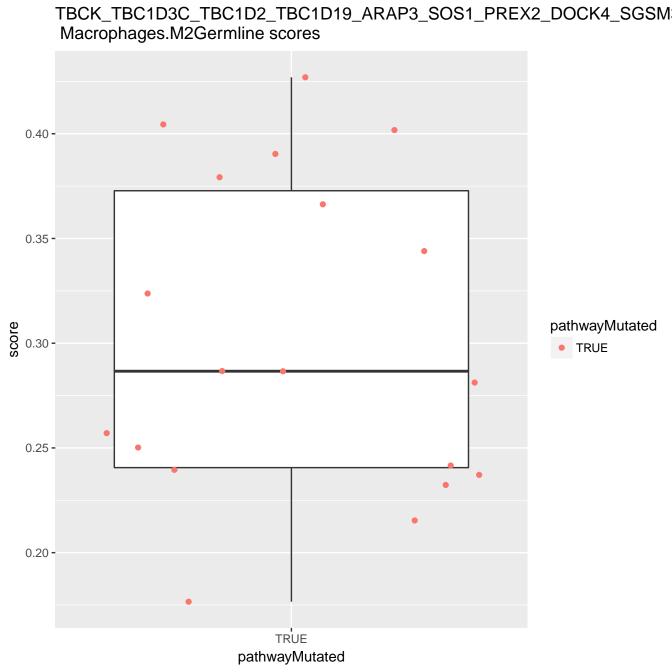
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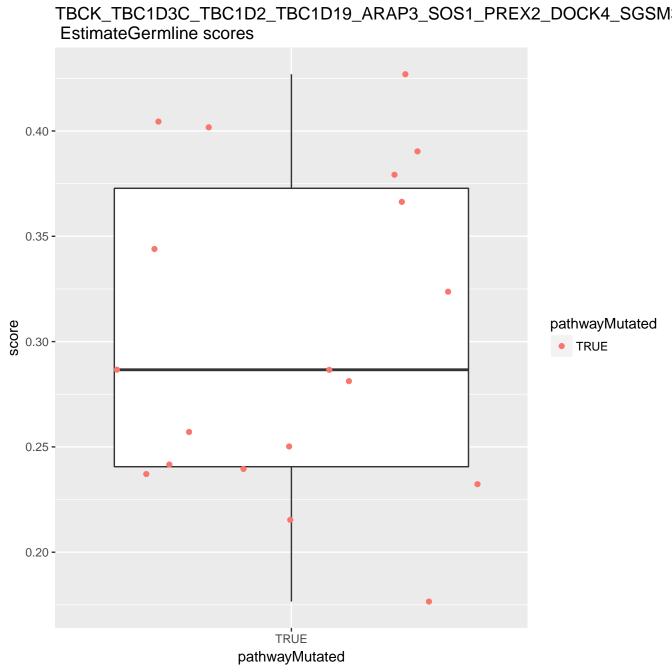


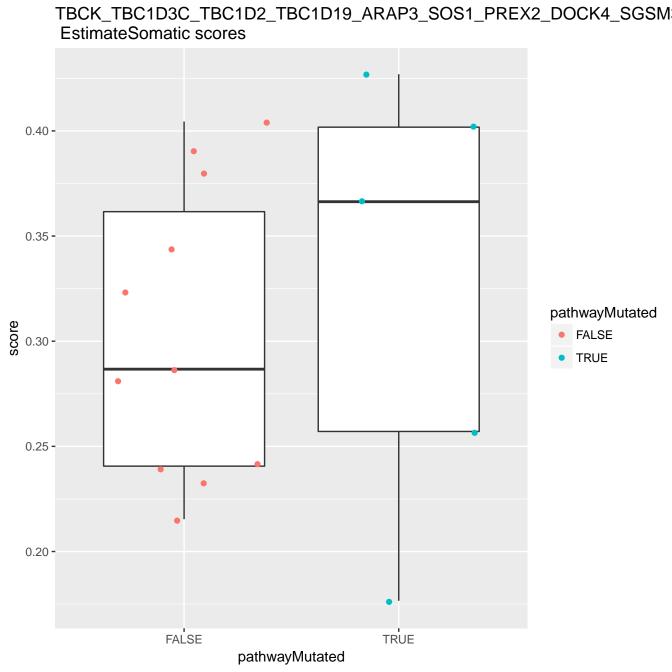


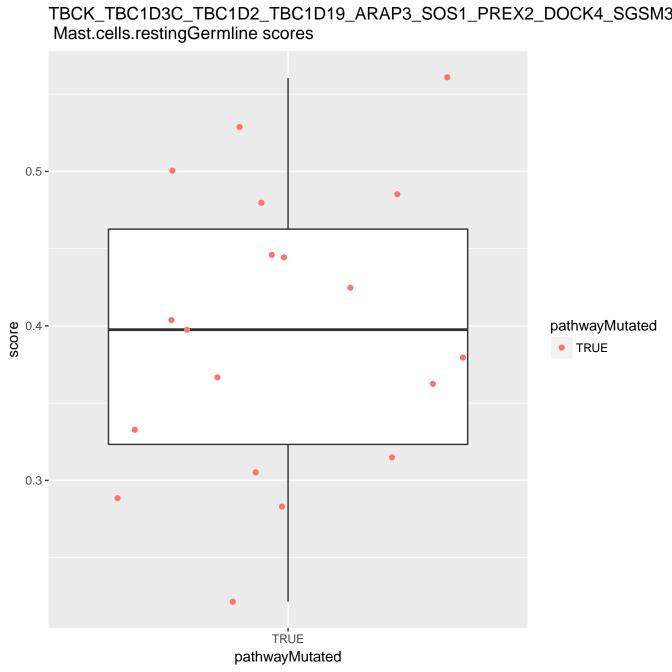
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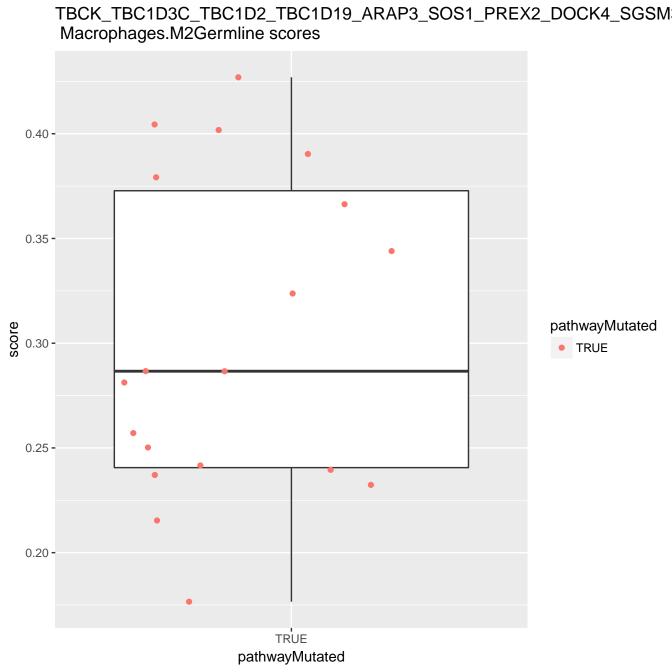
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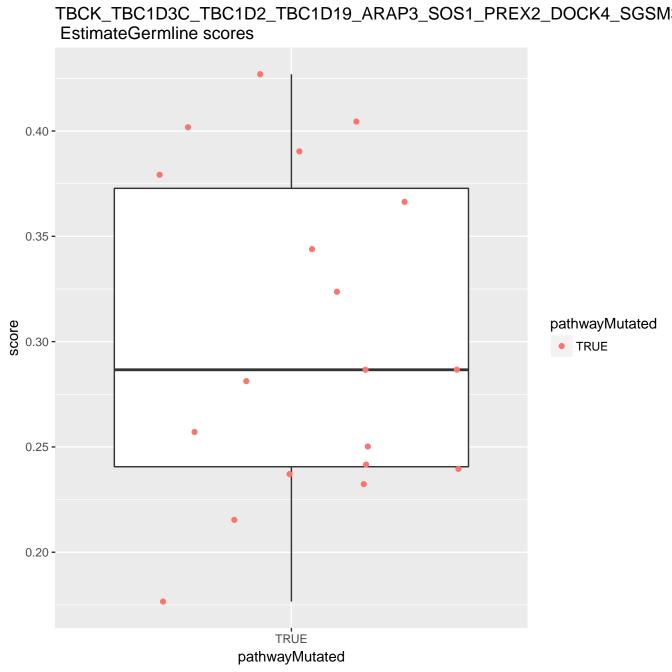


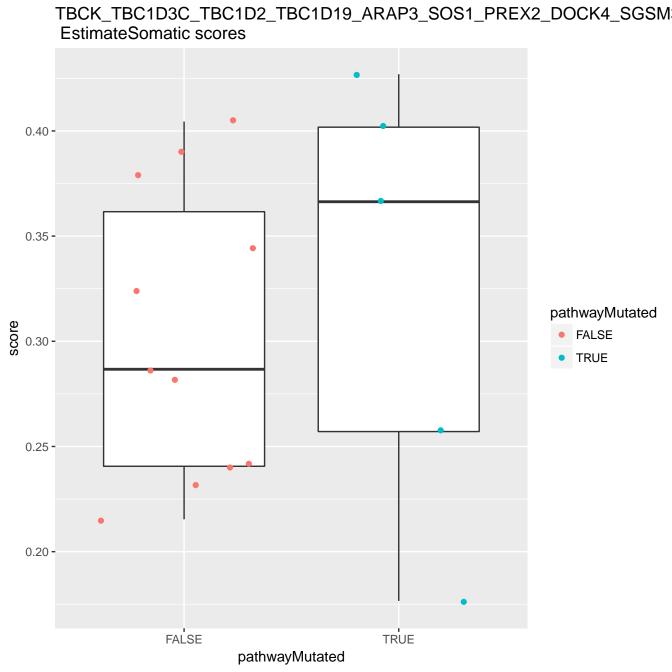


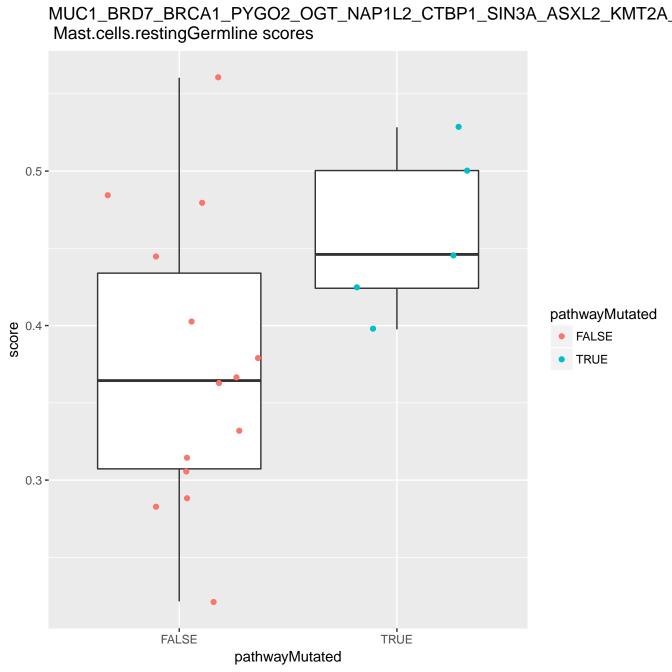
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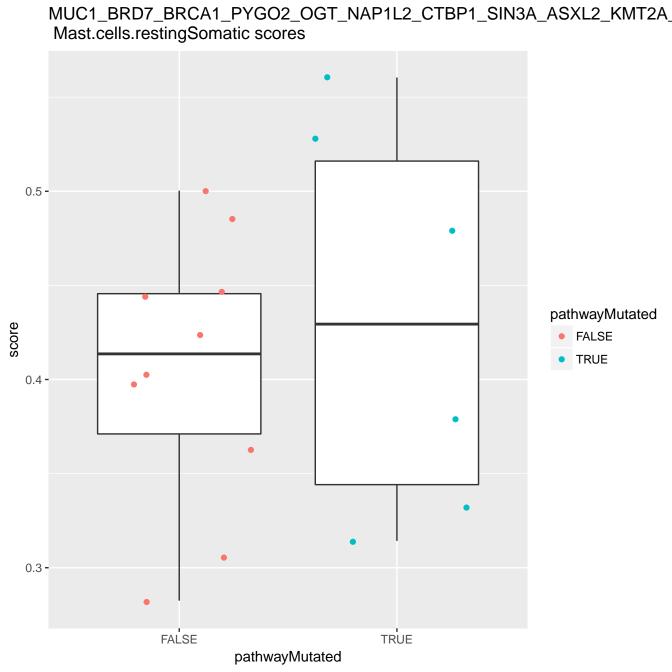


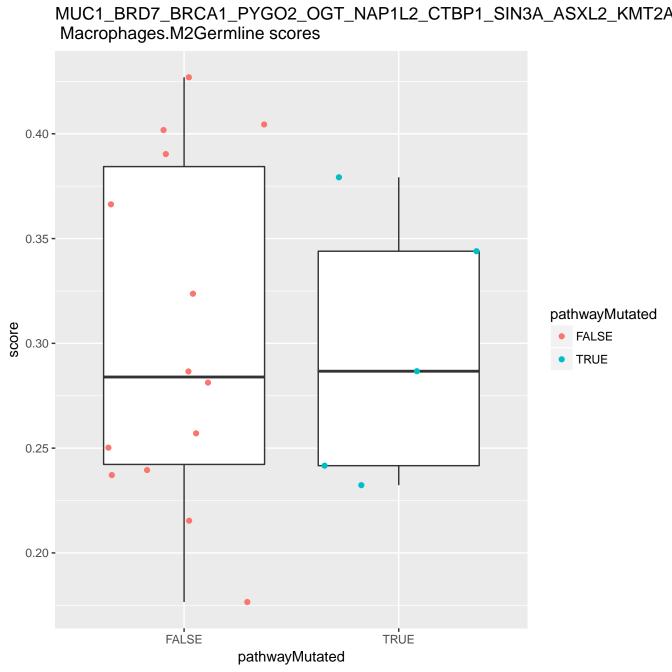
TBCK\_TBC1D3C\_TBC1D2\_TBC1D19\_ARAP3\_SOS1\_PREX2\_DOCK4\_SGSM Macrophages.M2Somatic scores 0.40 -0.35 pathwayMutated **SCOre** 0.30 -FALSE TRUE 0.25 -0.20 -FALSE TRUE pathwayMutated

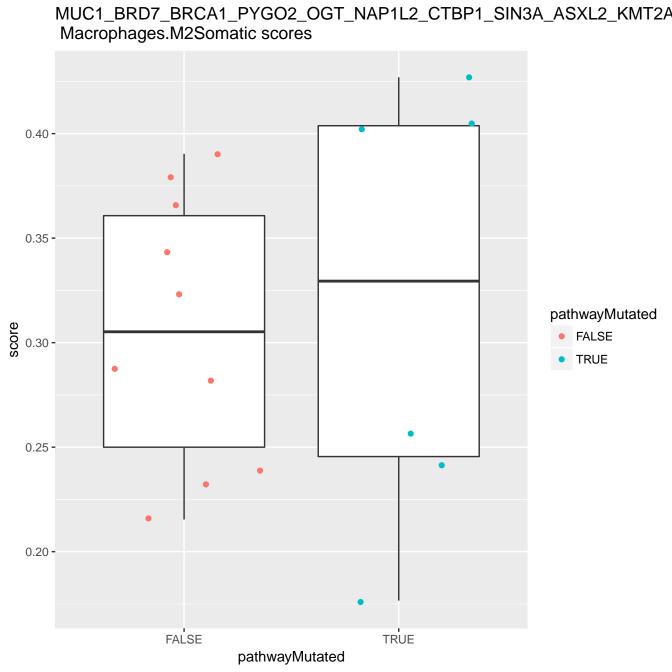


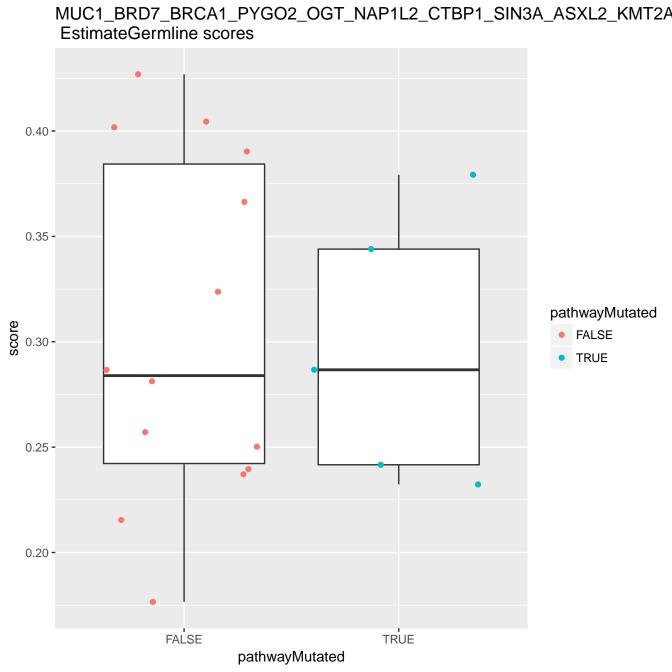


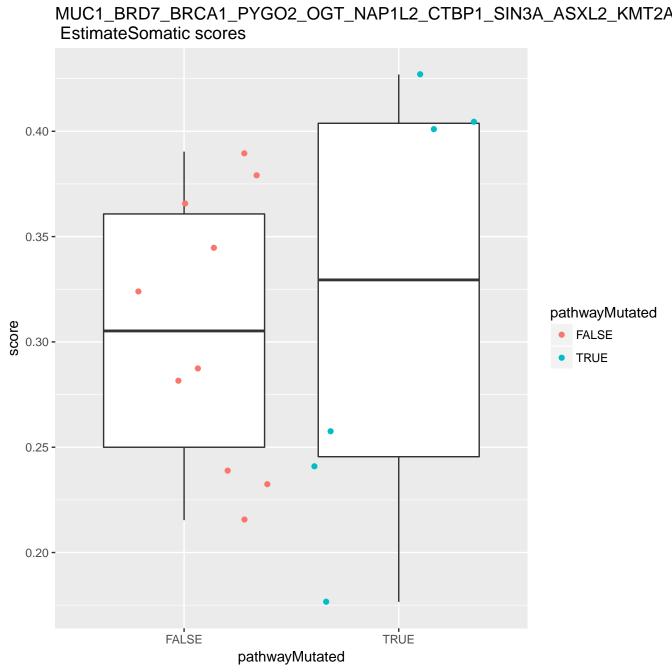


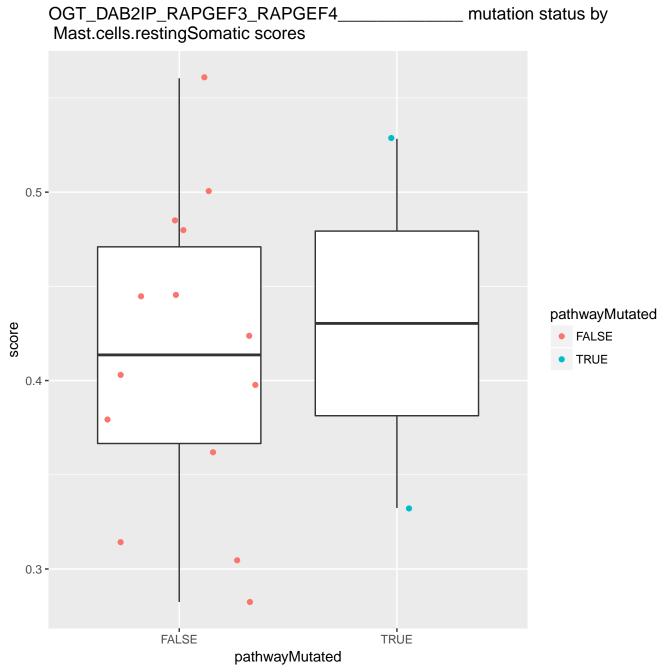


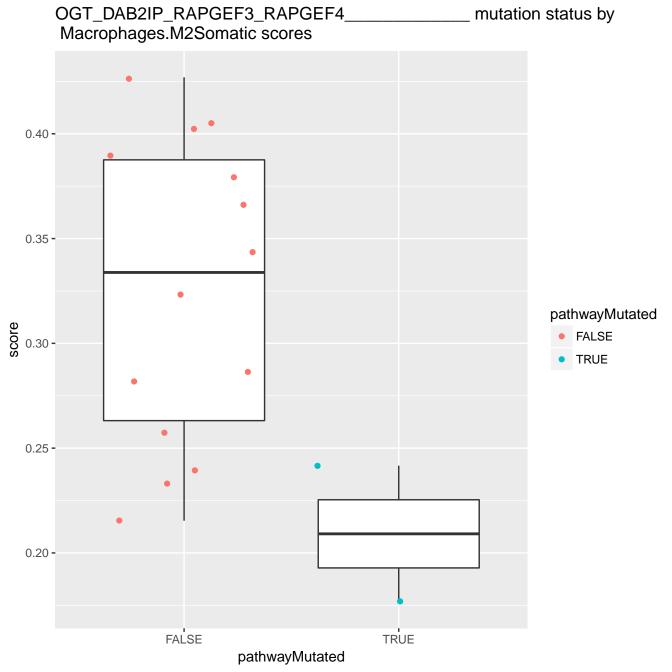


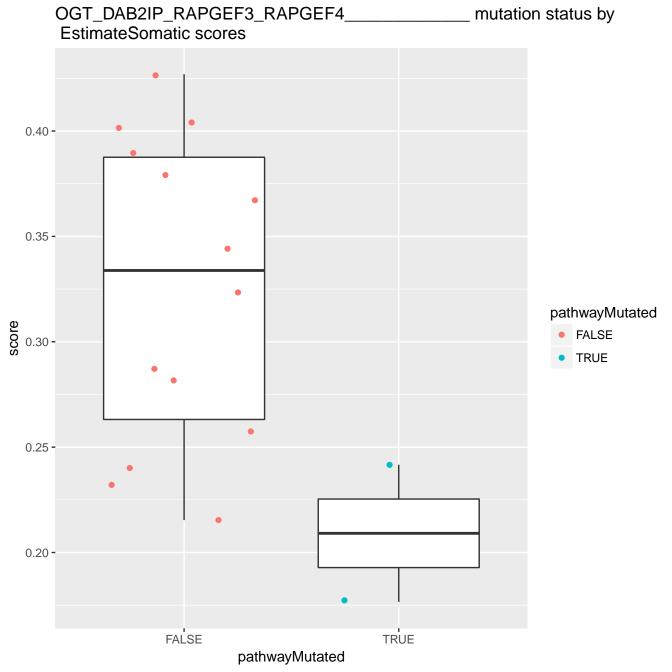


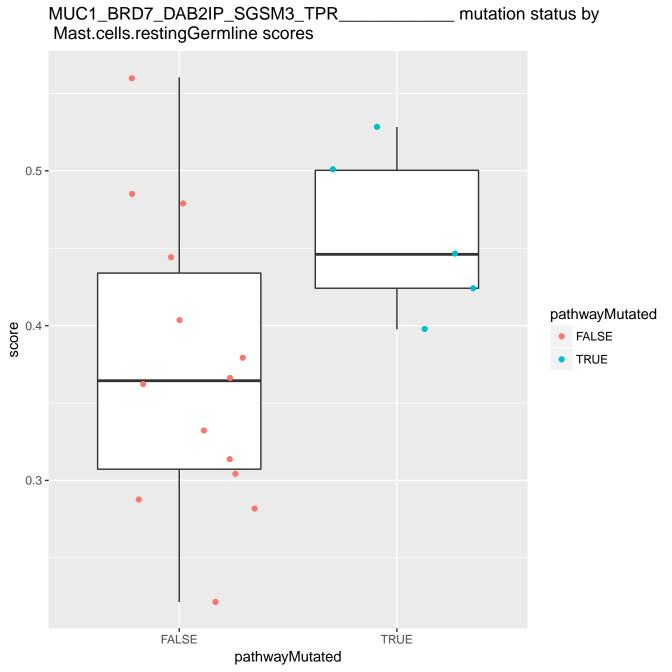


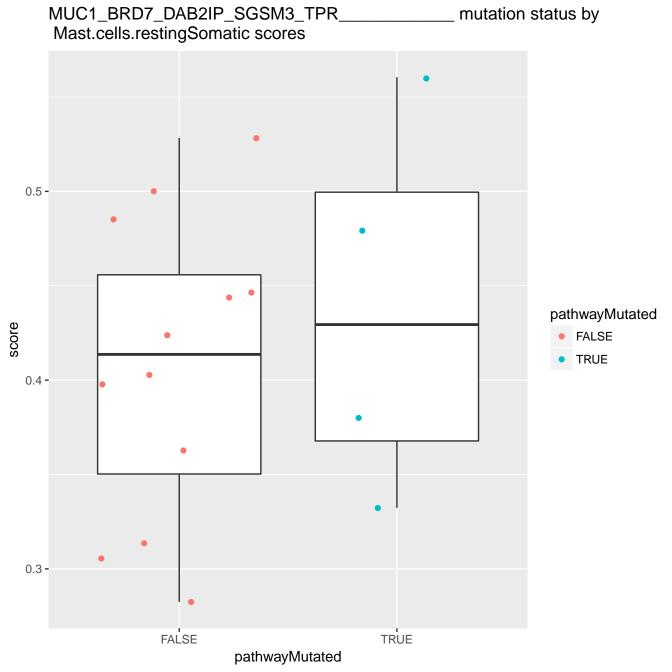


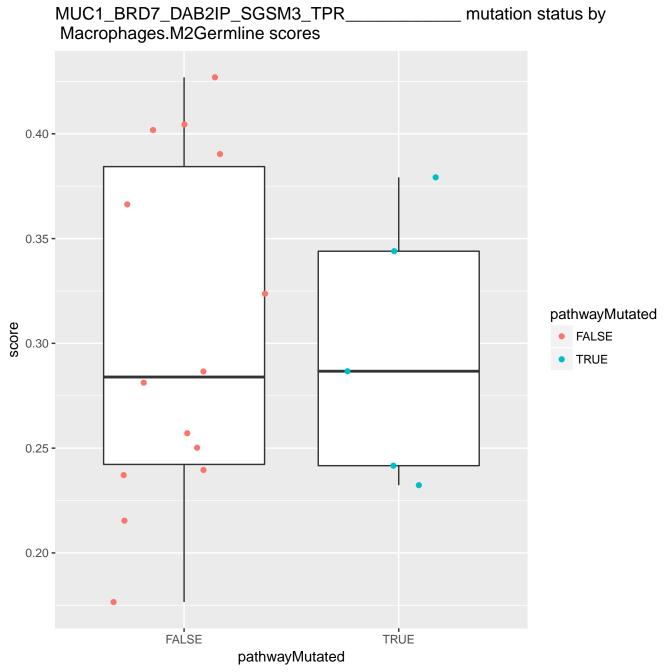


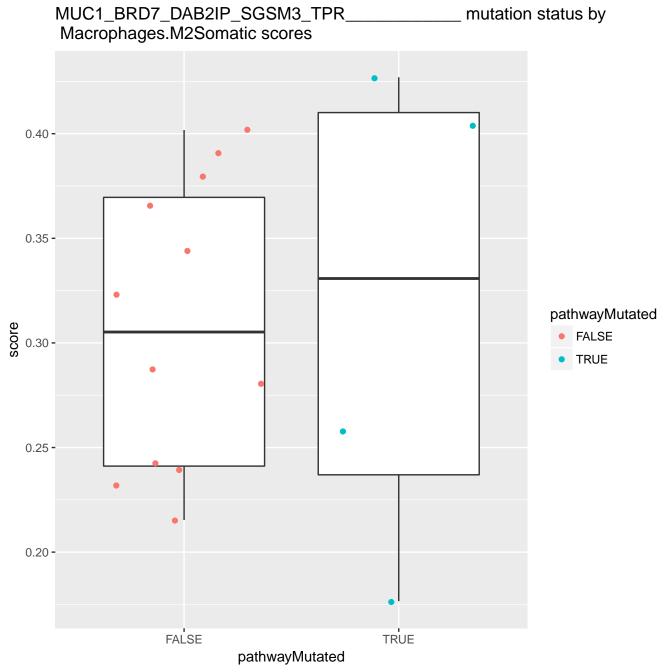


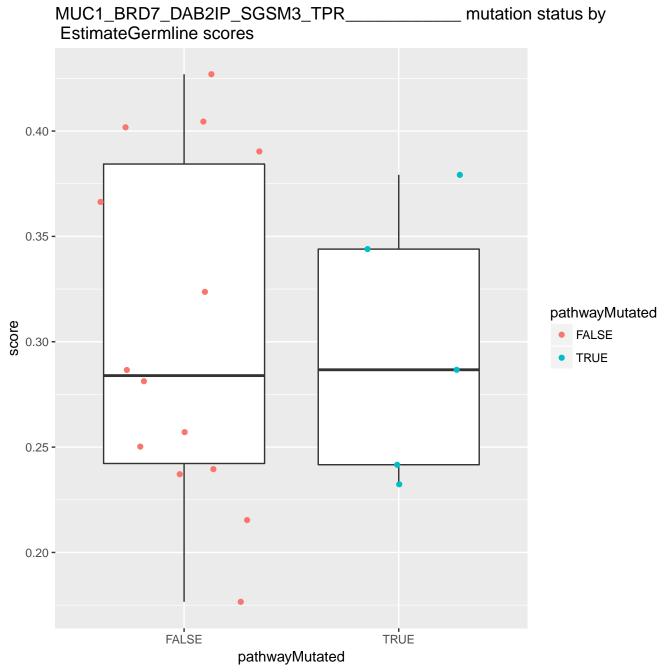


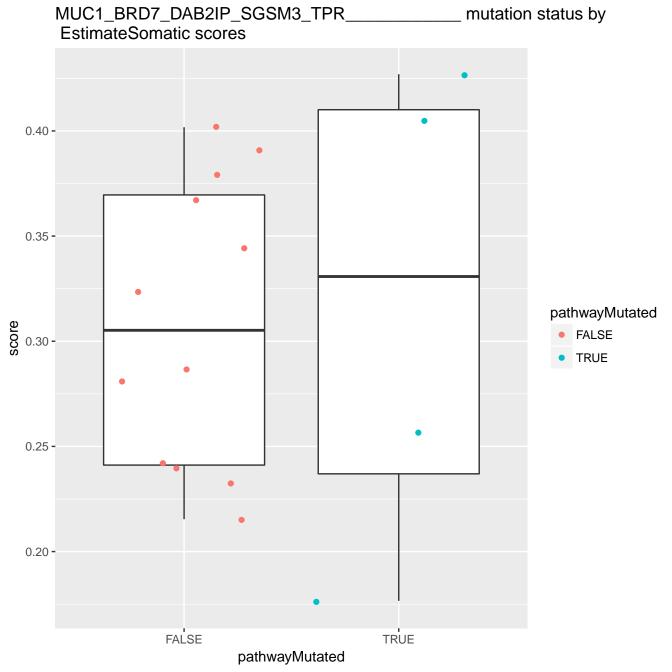


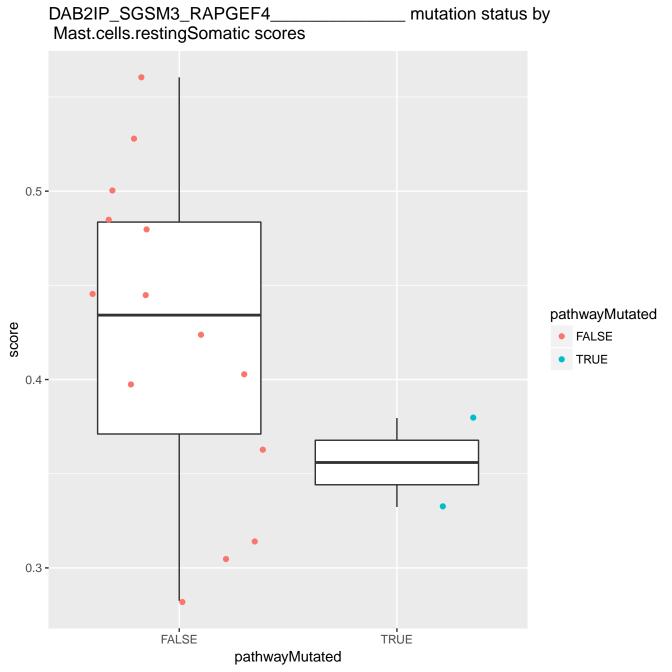


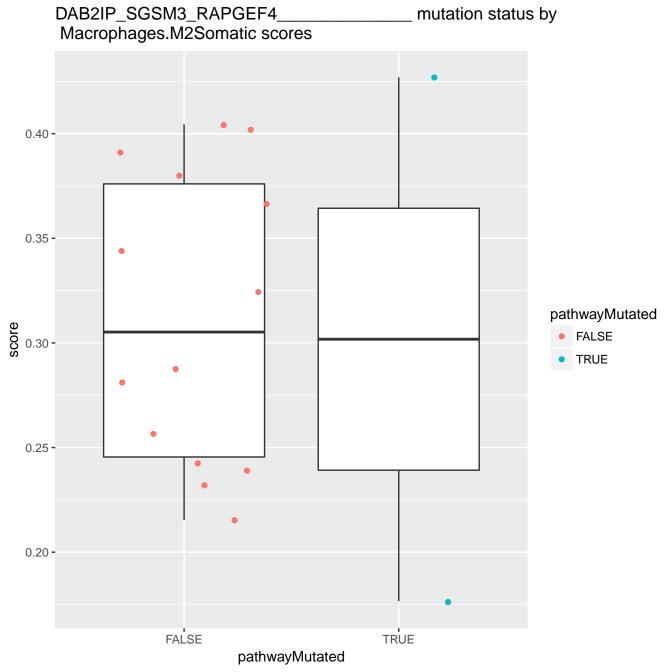


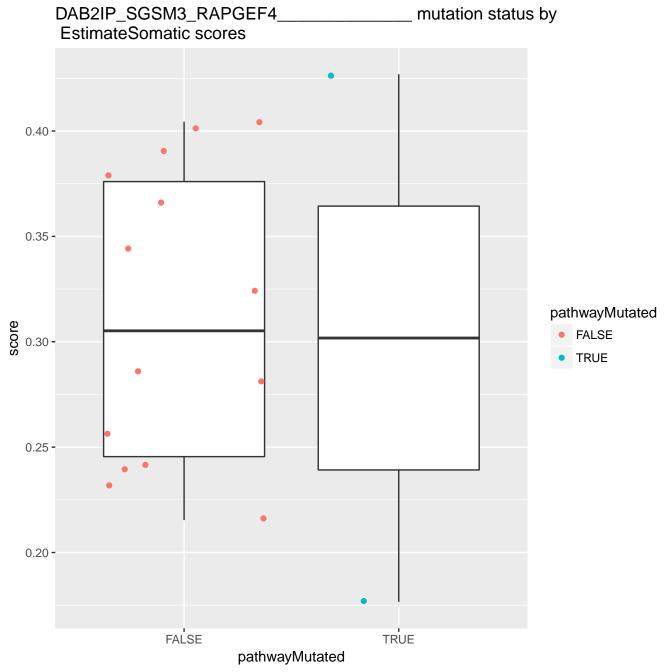


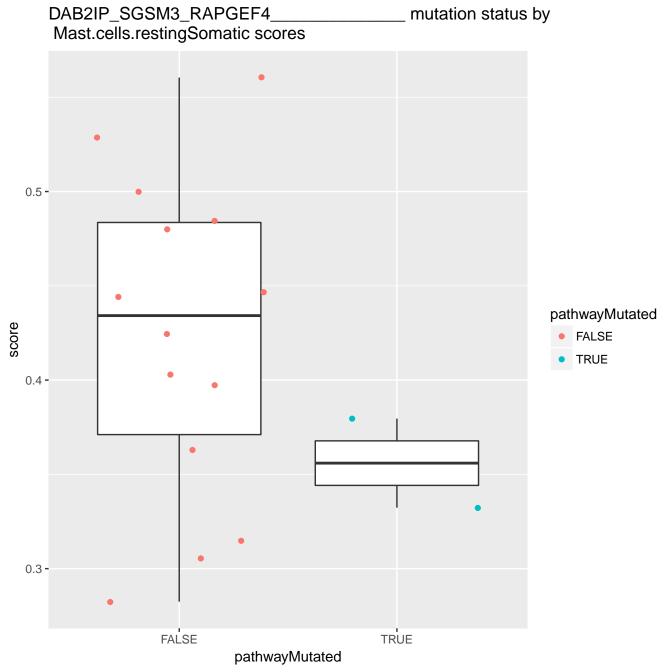


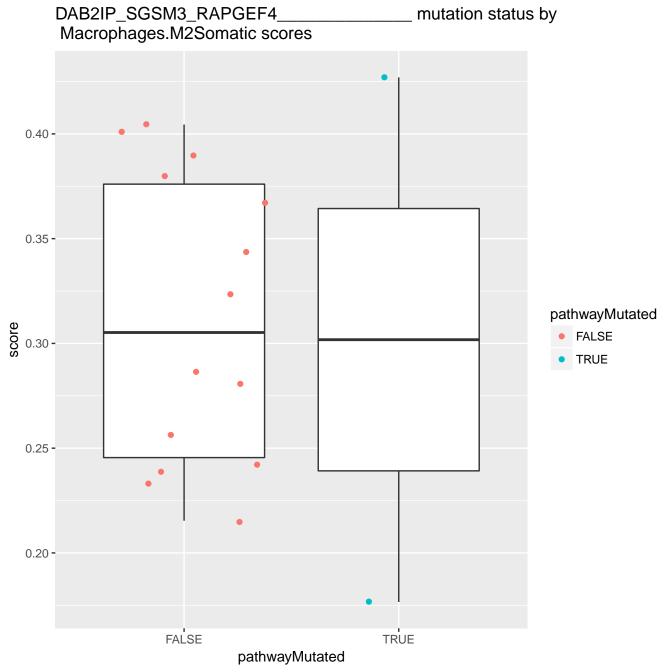


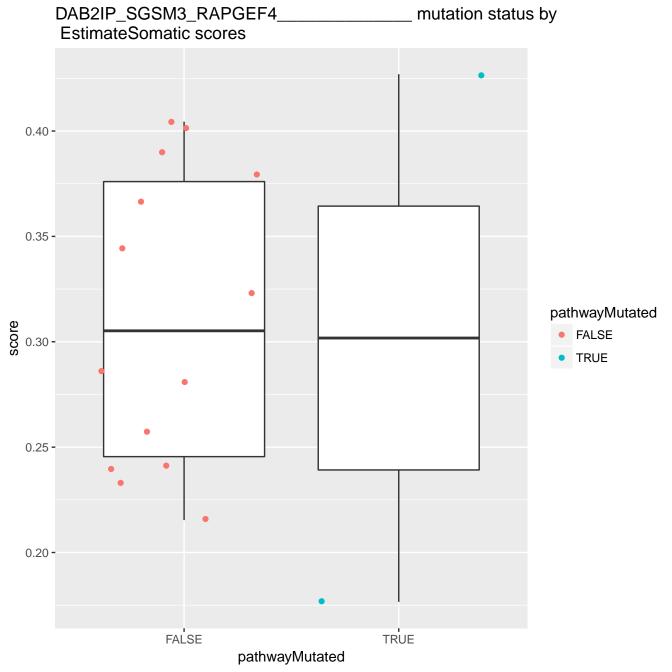


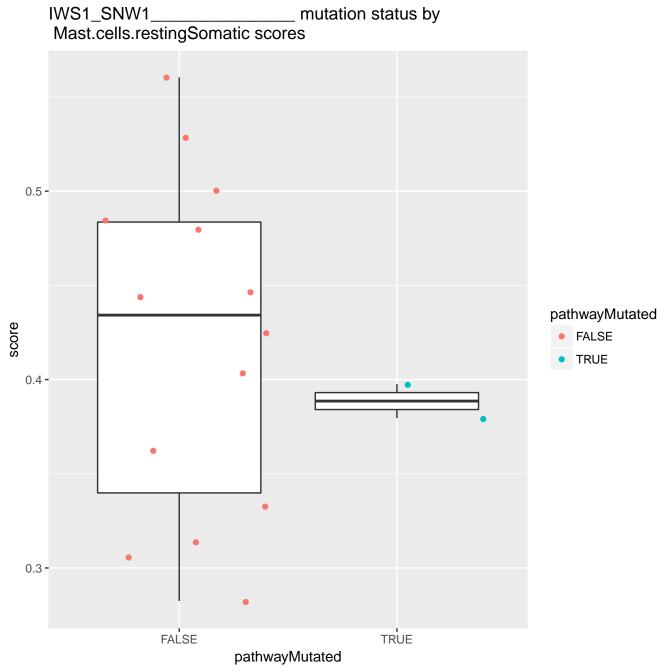


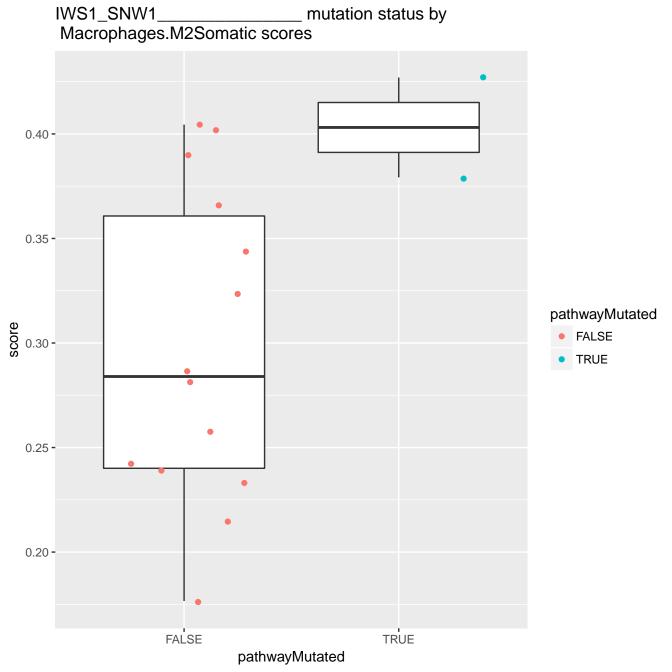


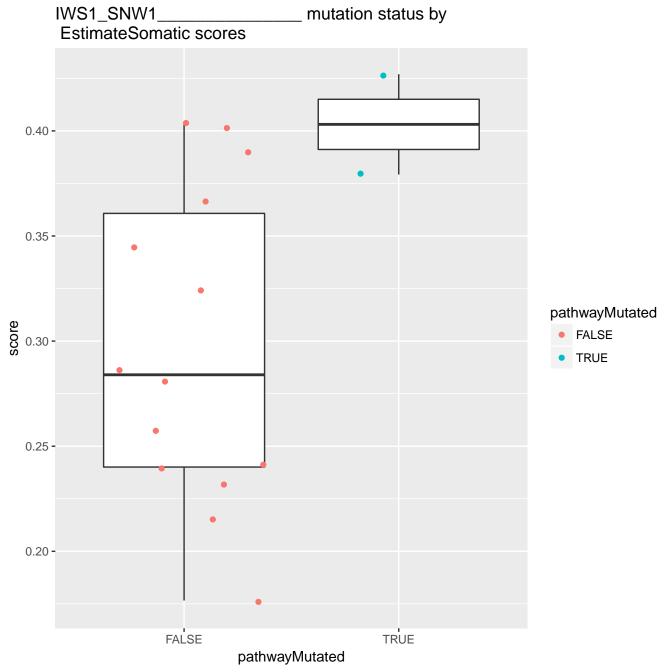


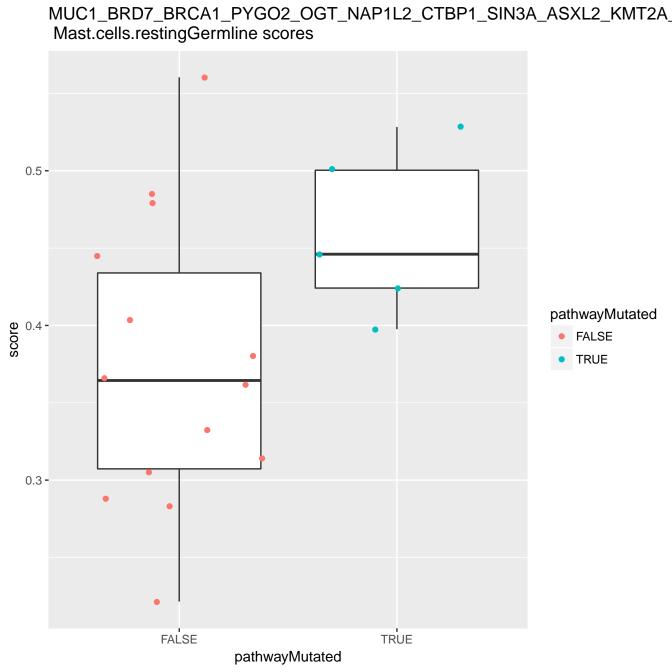


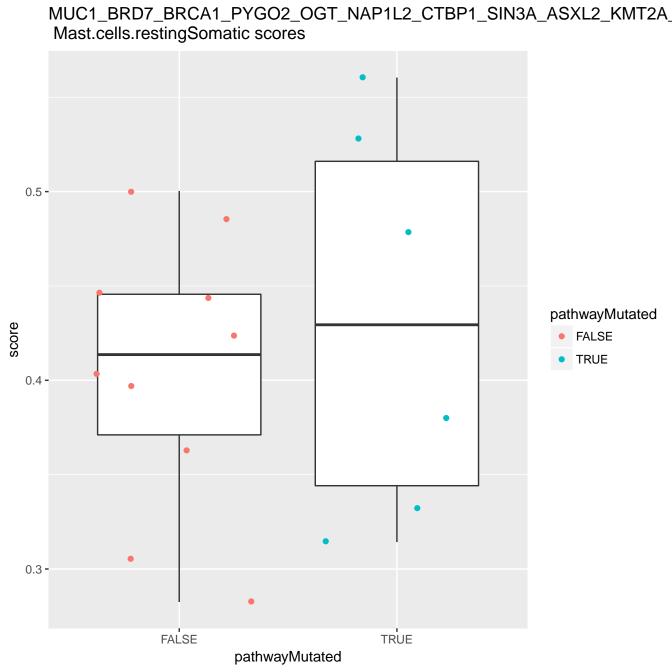


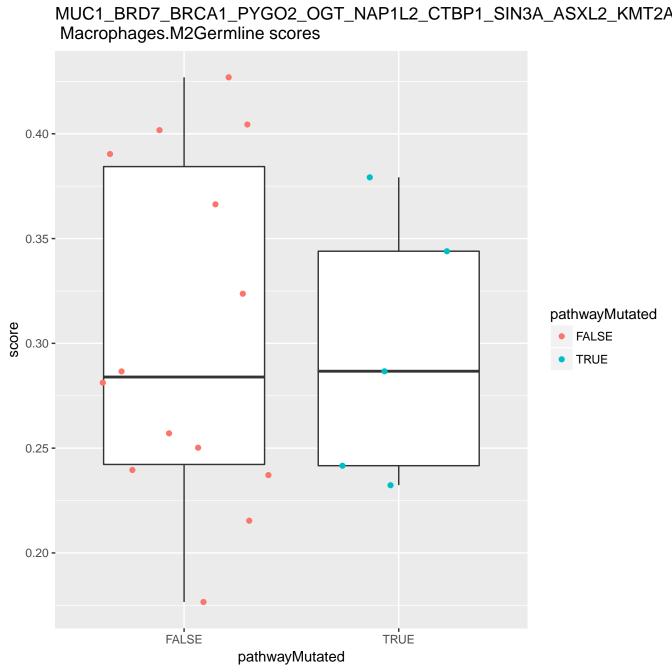


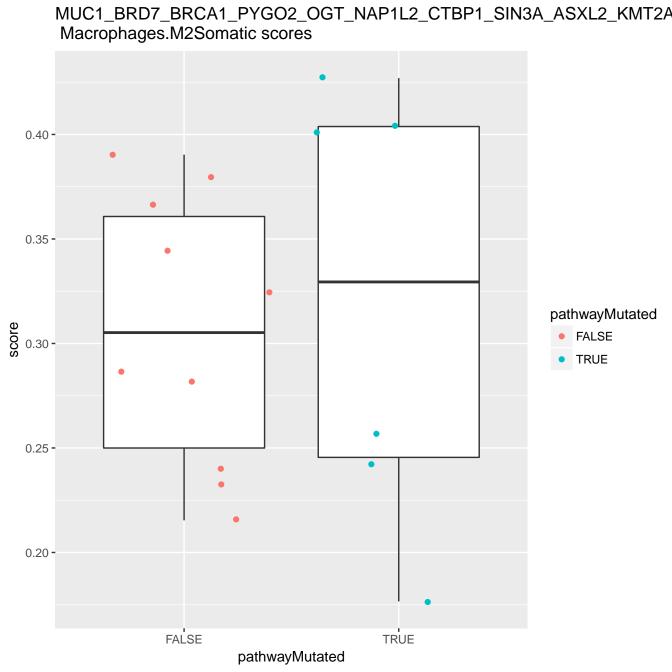


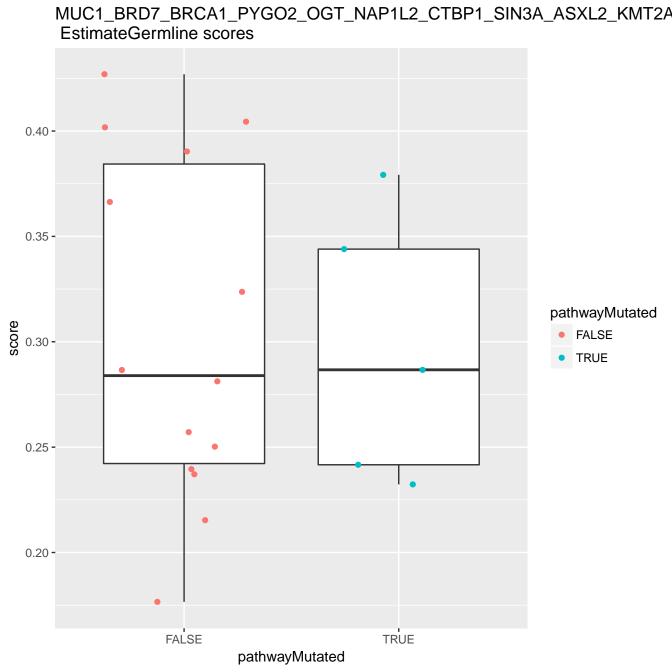


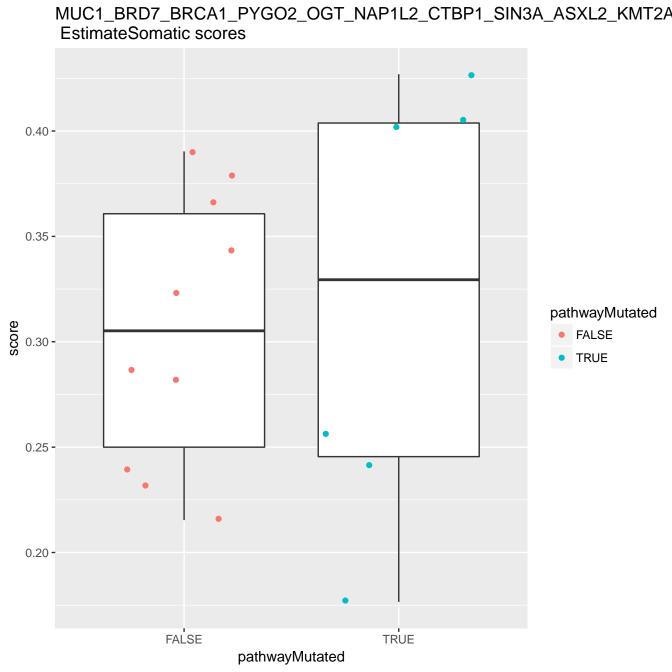


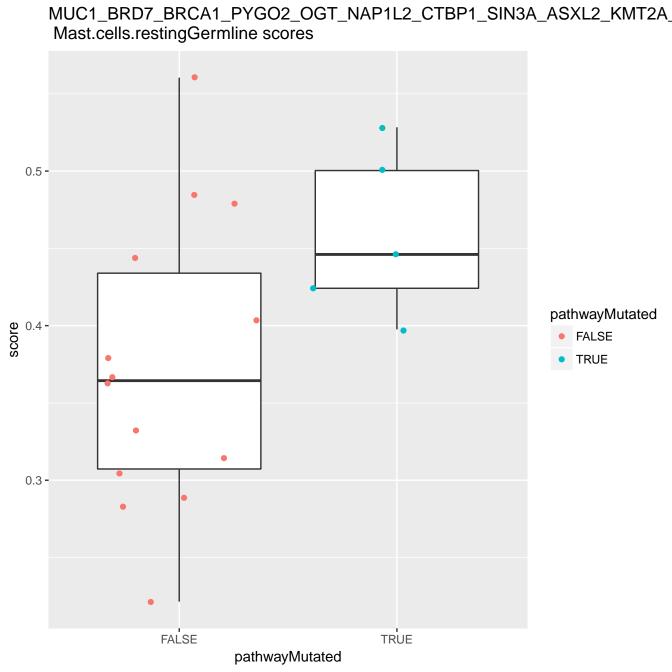


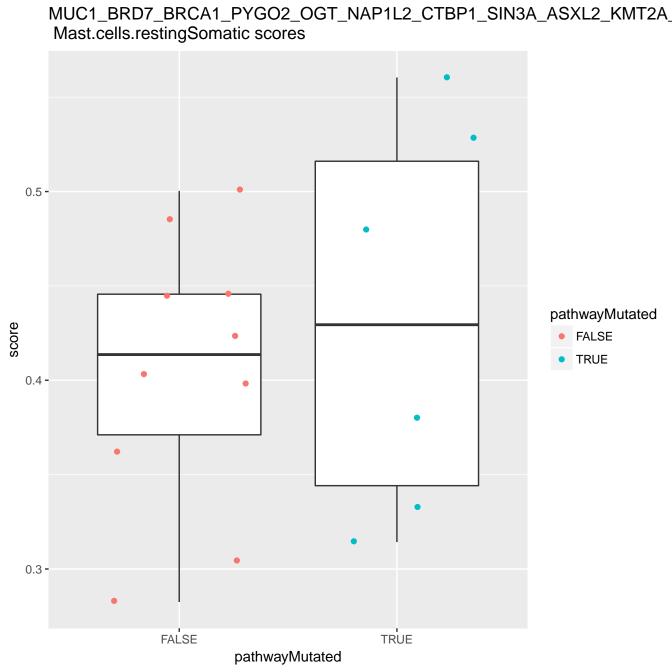


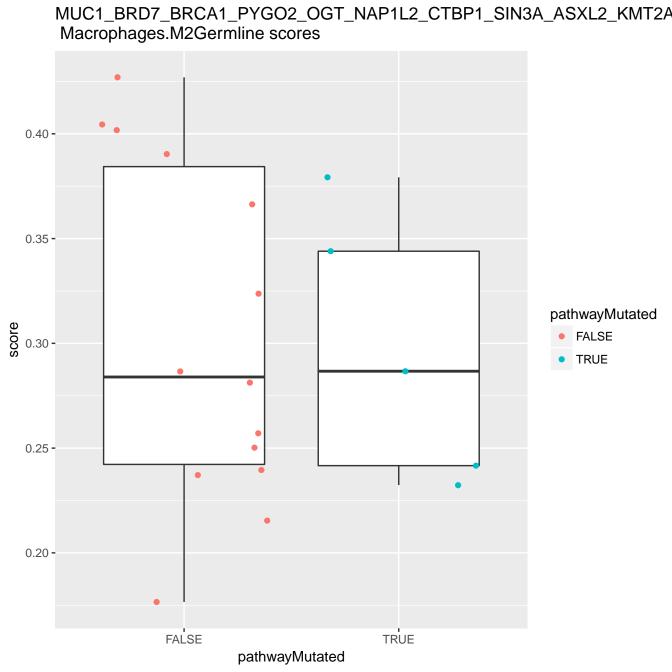


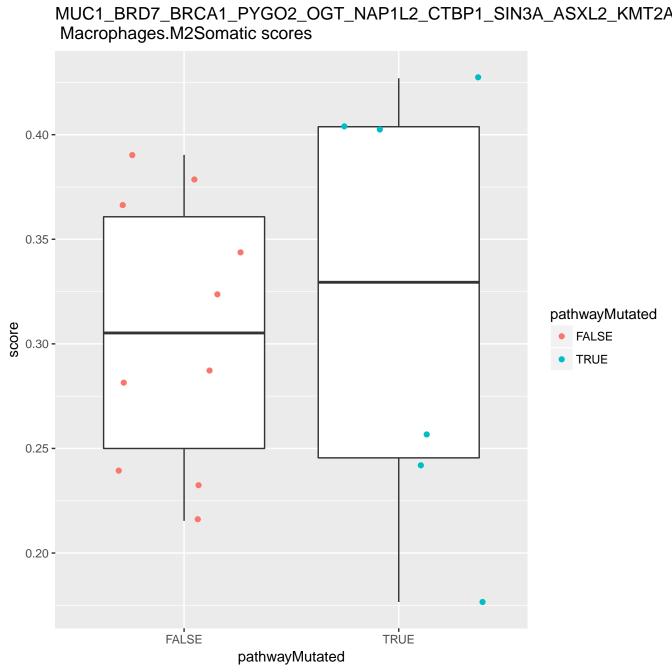


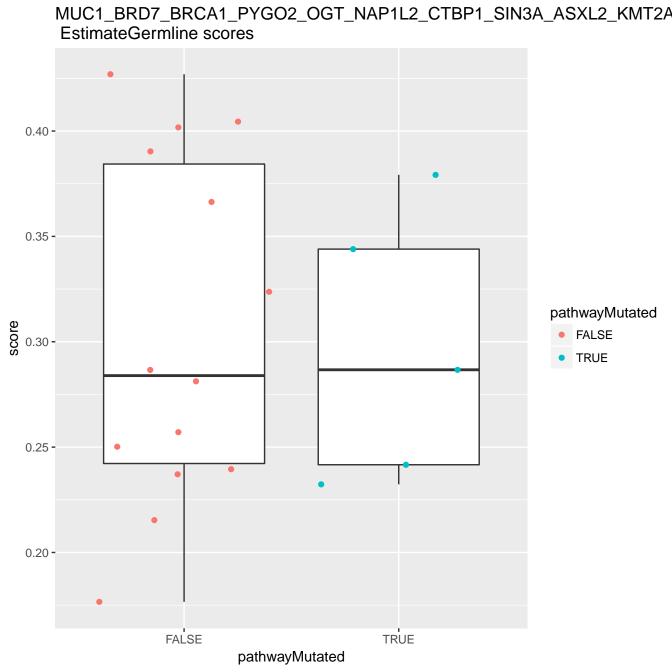


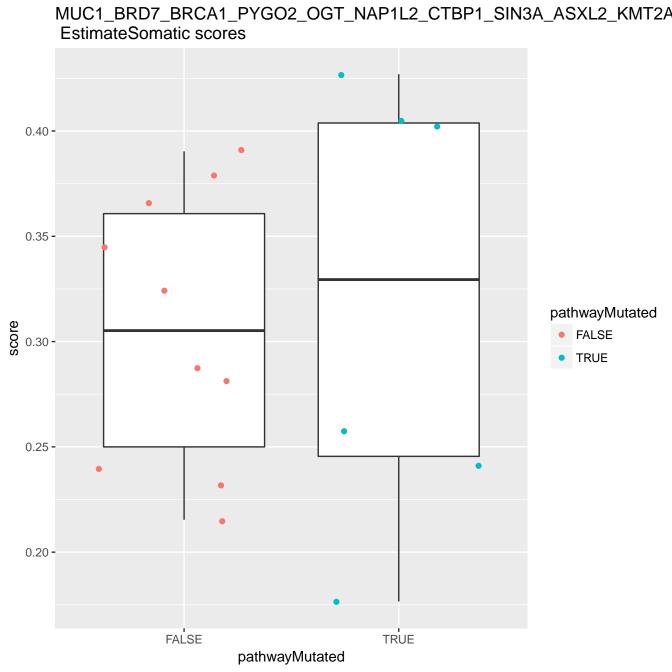


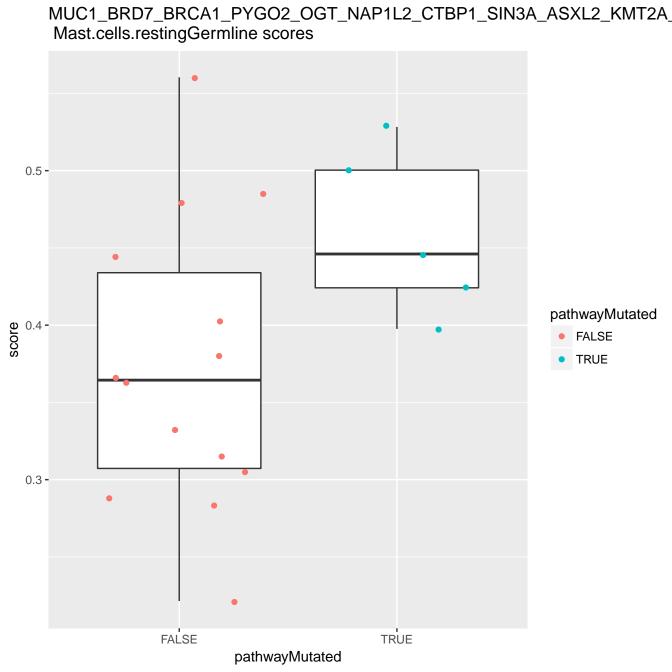


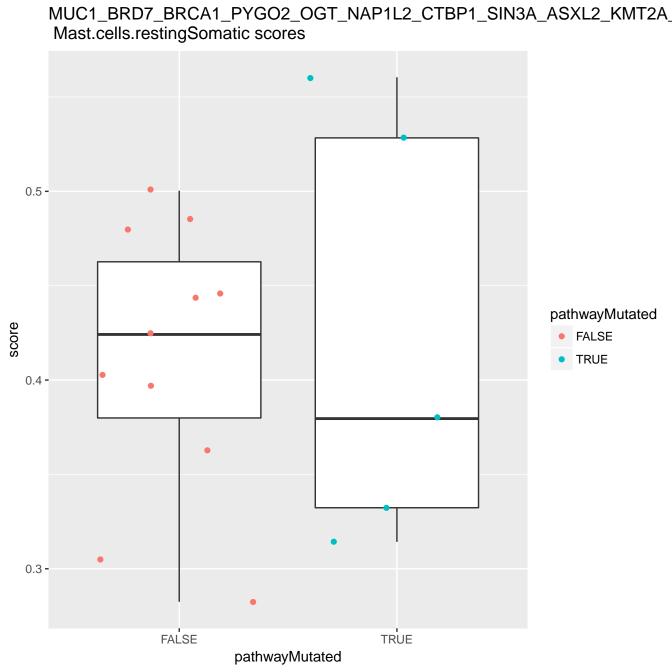


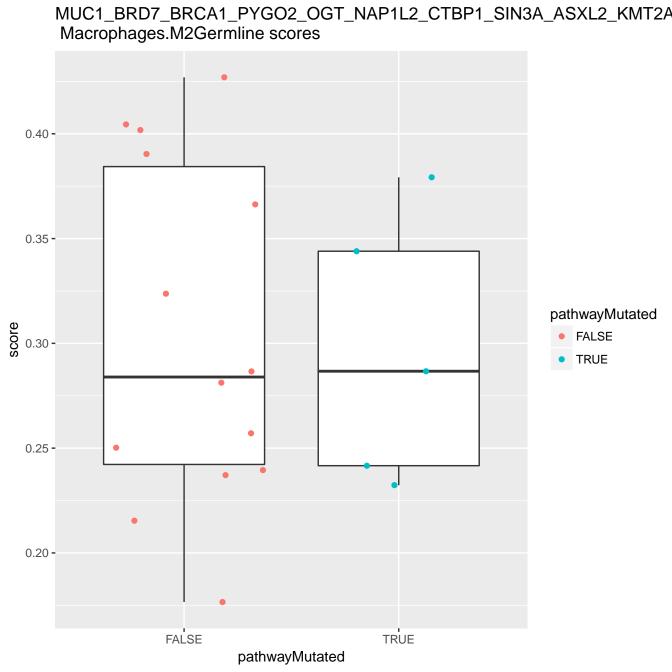


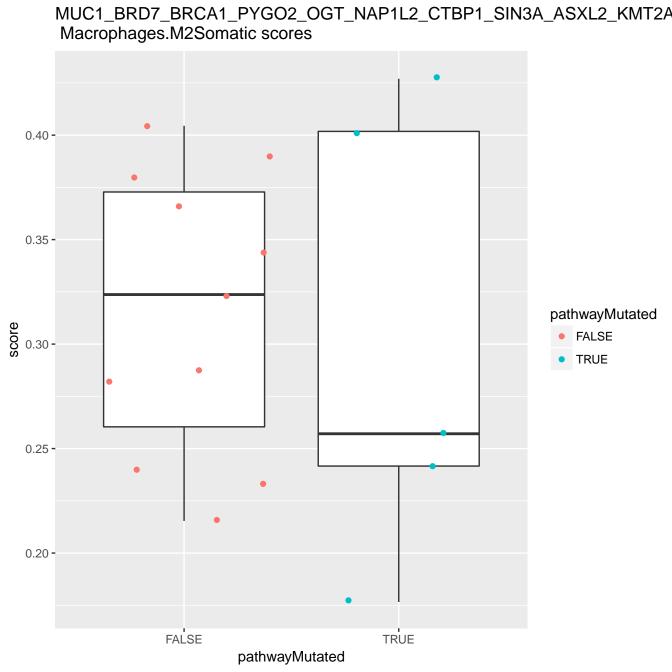


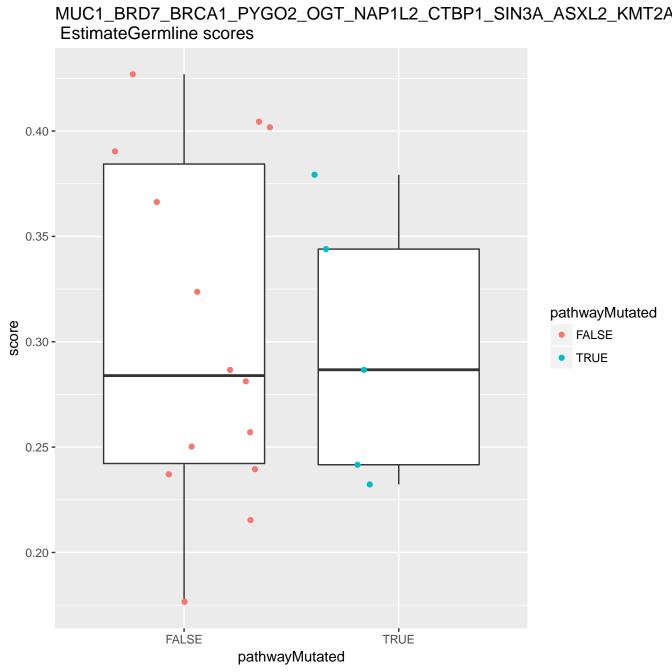


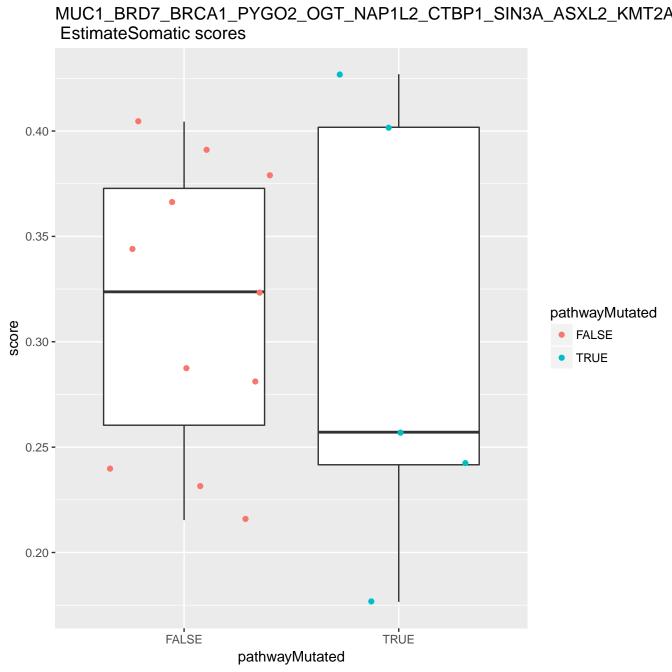


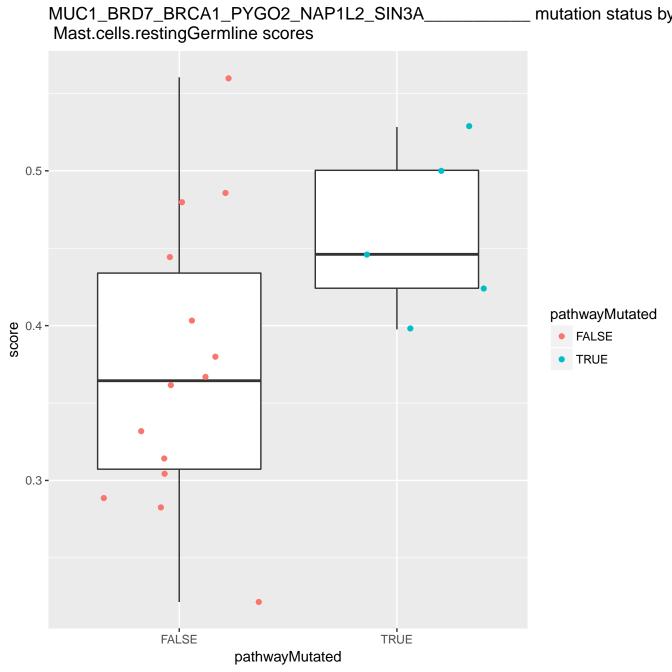


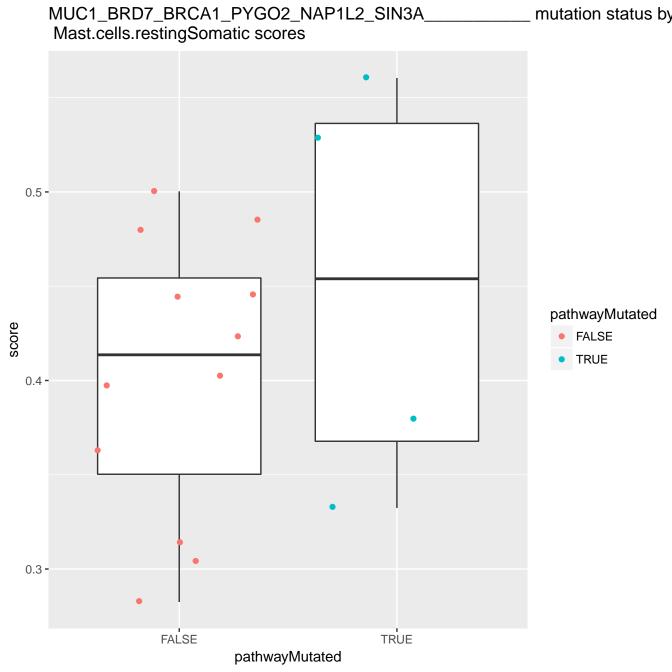


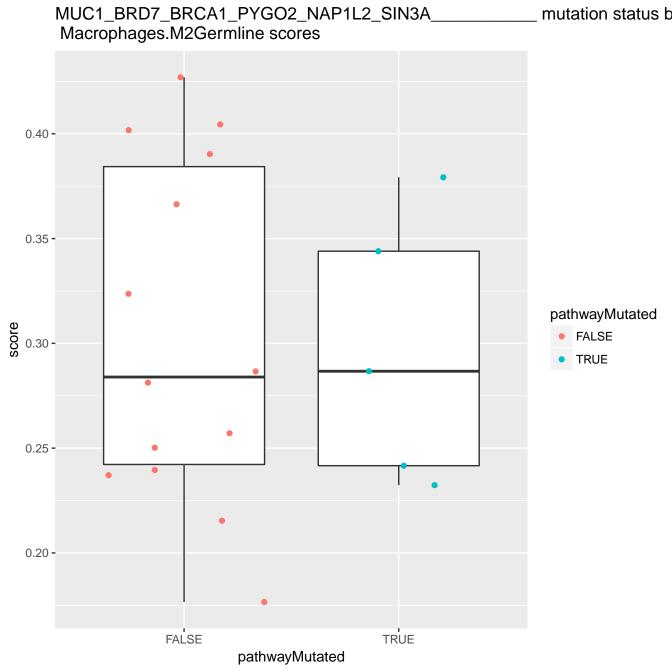


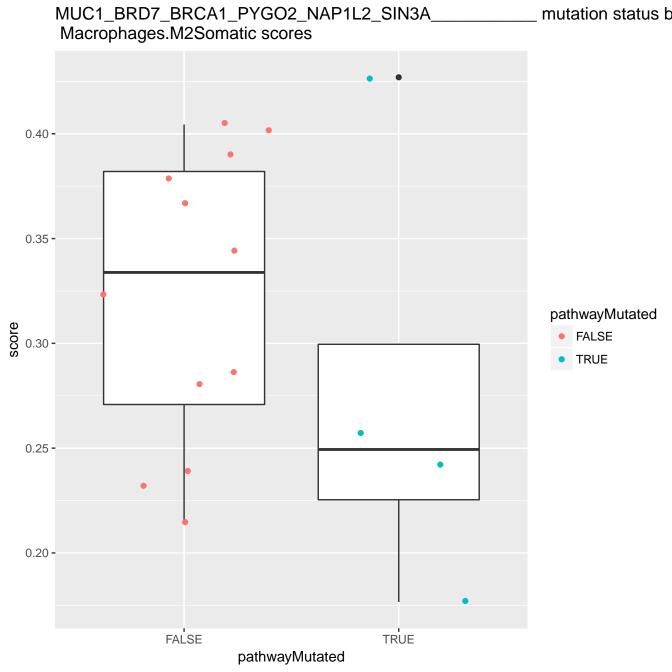


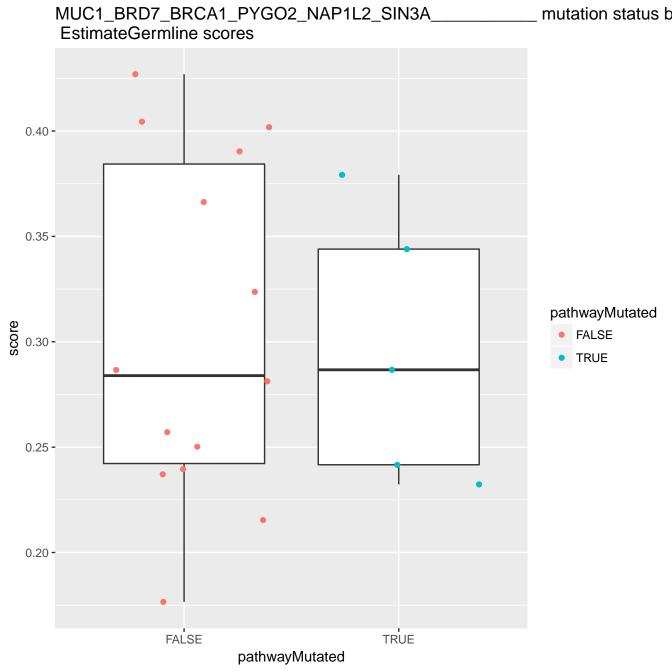


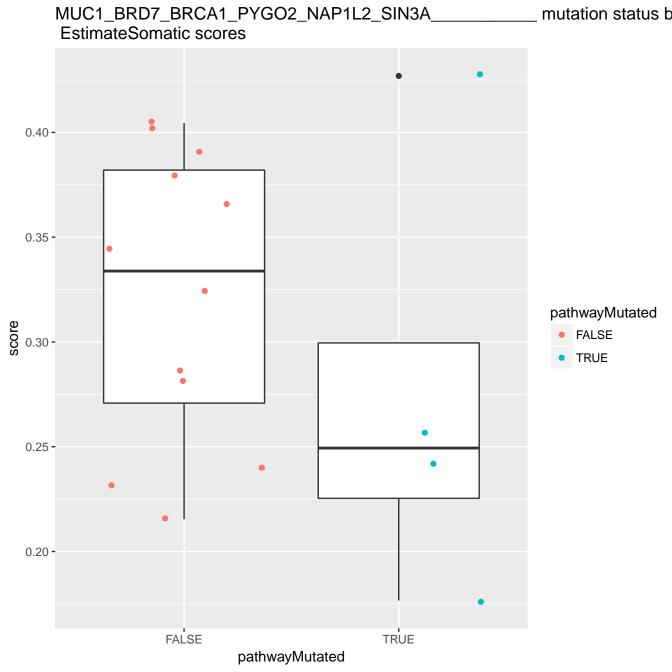


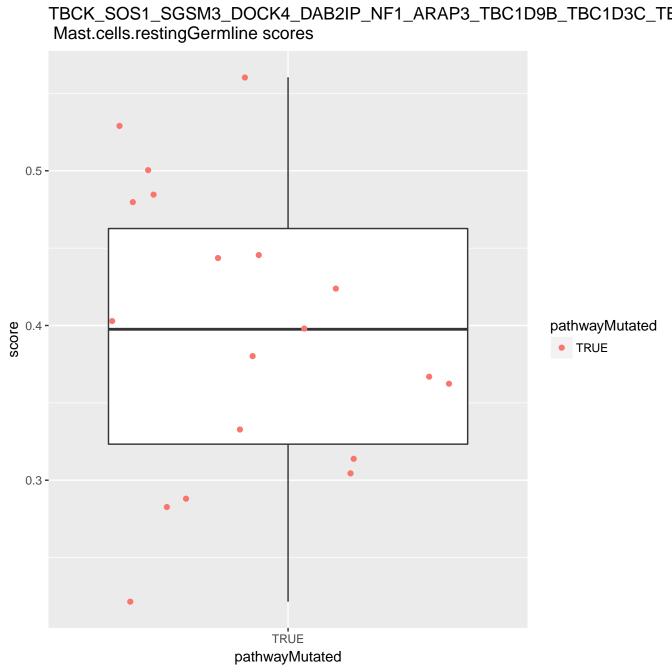


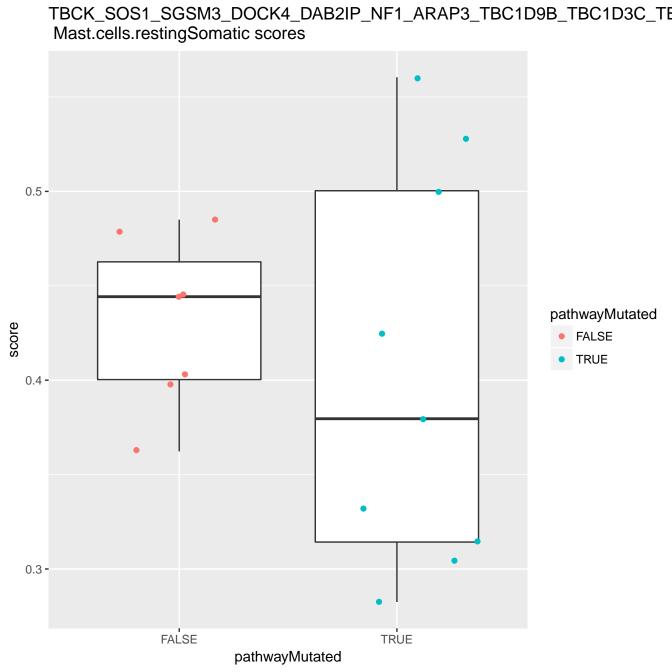


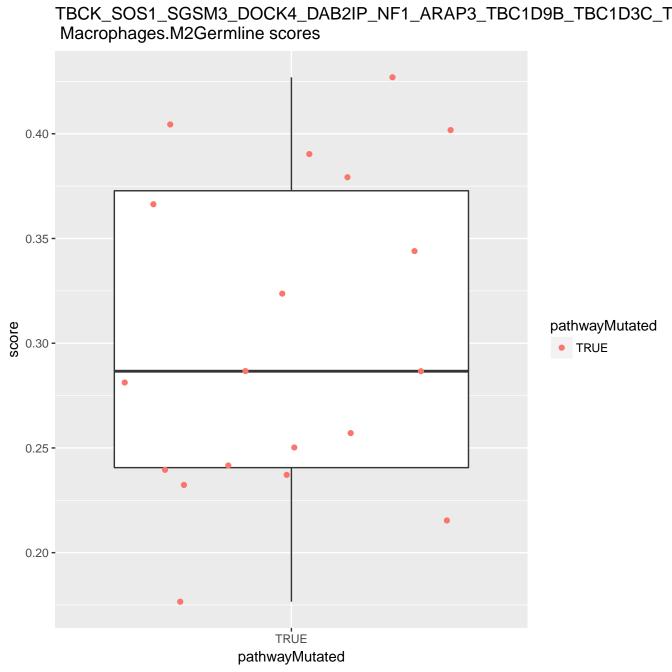




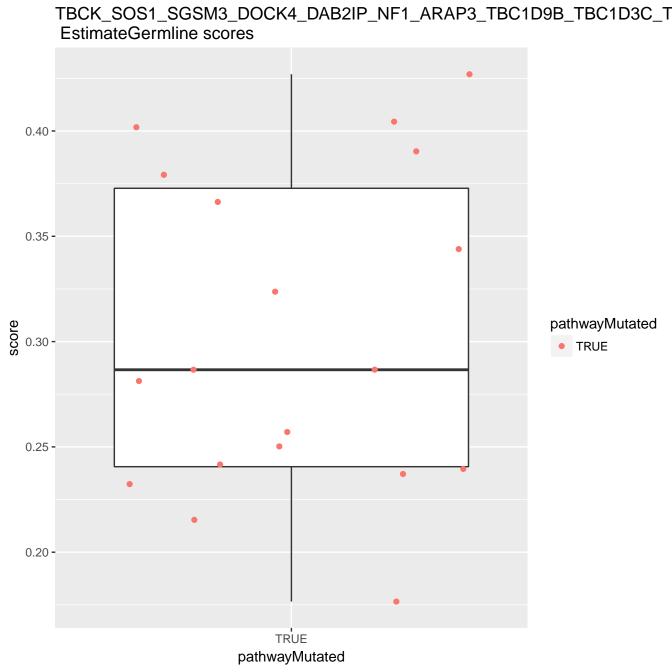


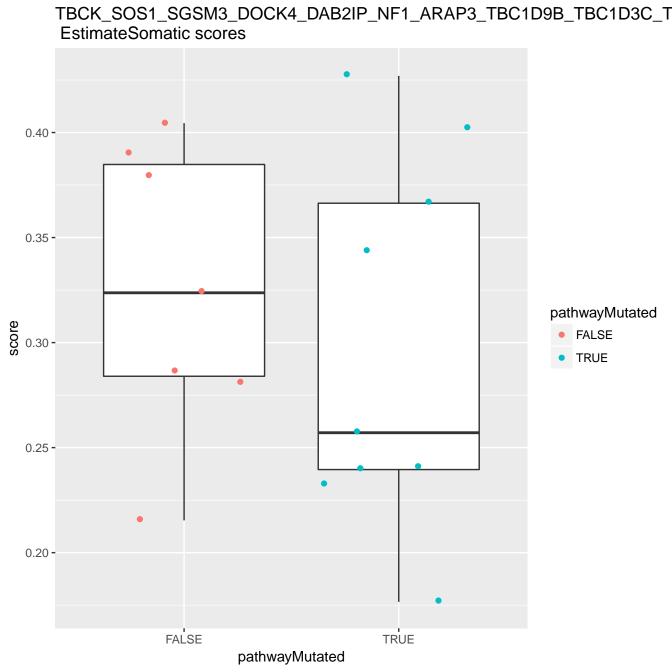


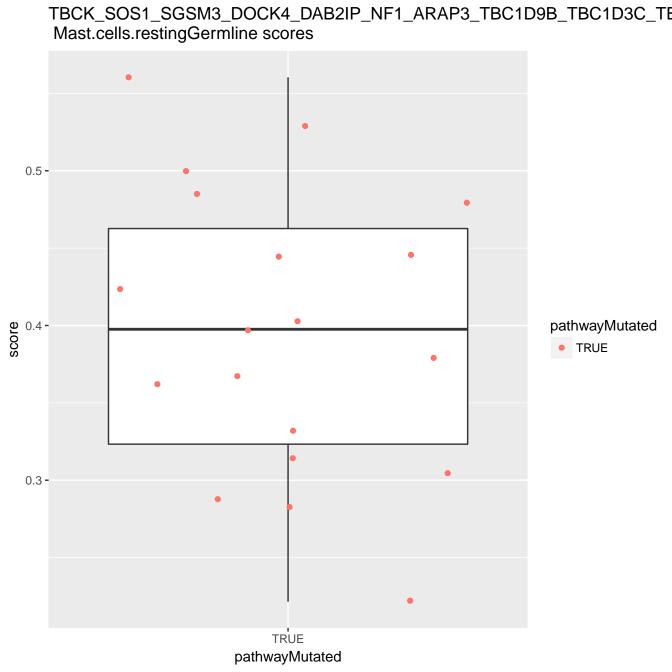


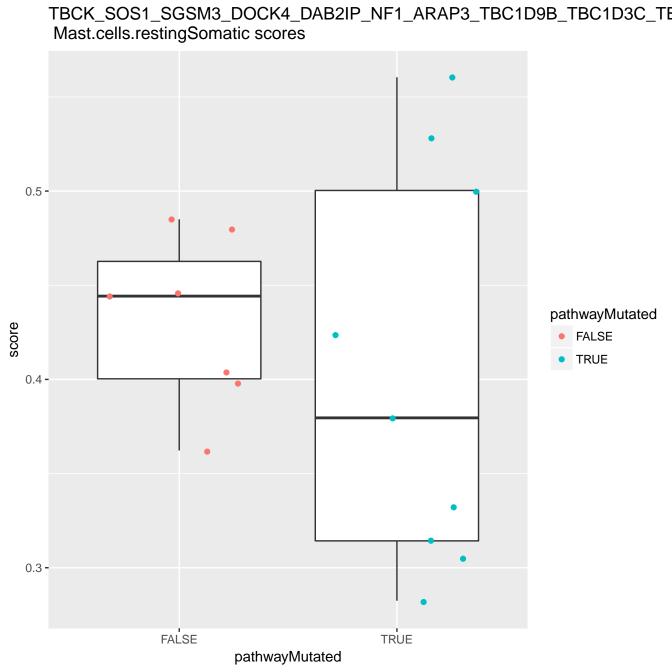


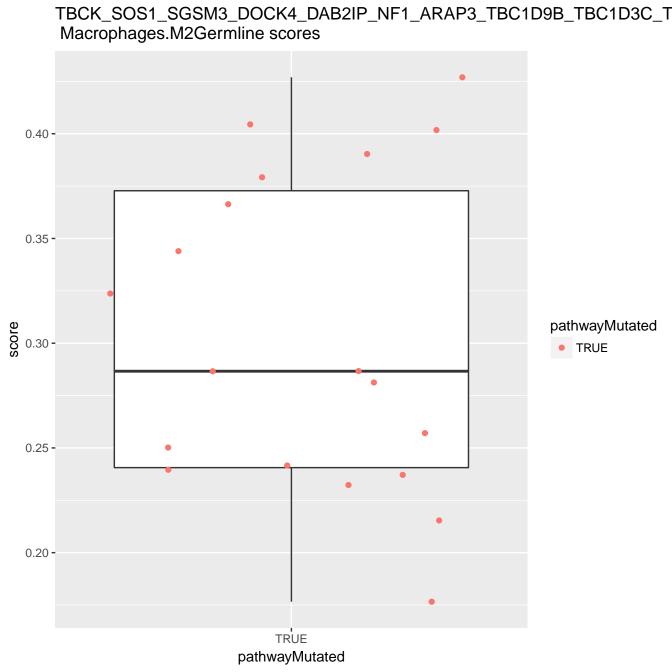
TBCK\_SOS1\_SGSM3\_DOCK4\_DAB2IP\_NF1\_ARAP3\_TBC1D9B\_TBC1D3C\_T Macrophages.M2Somatic scores 0.40 -0.35 pathwayMutated SCO 6 0.30 -FALSE TRUE 0.25 -0.20 -FALSE TRUE pathwayMutated



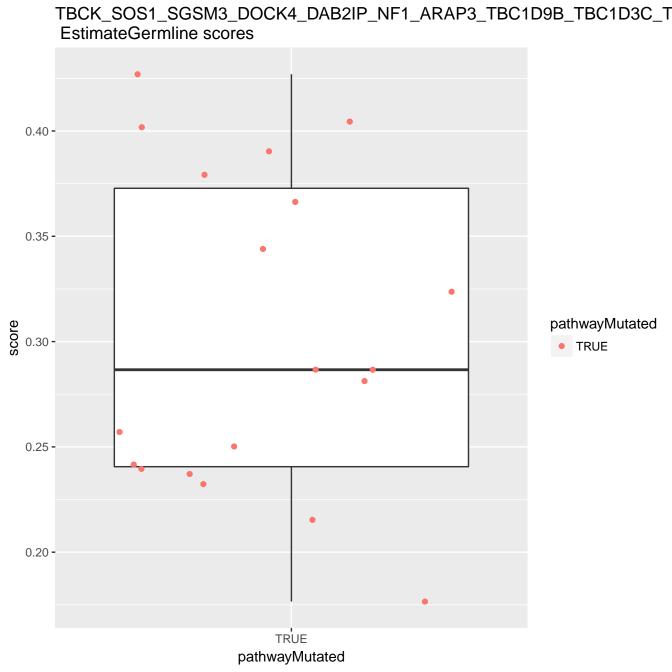


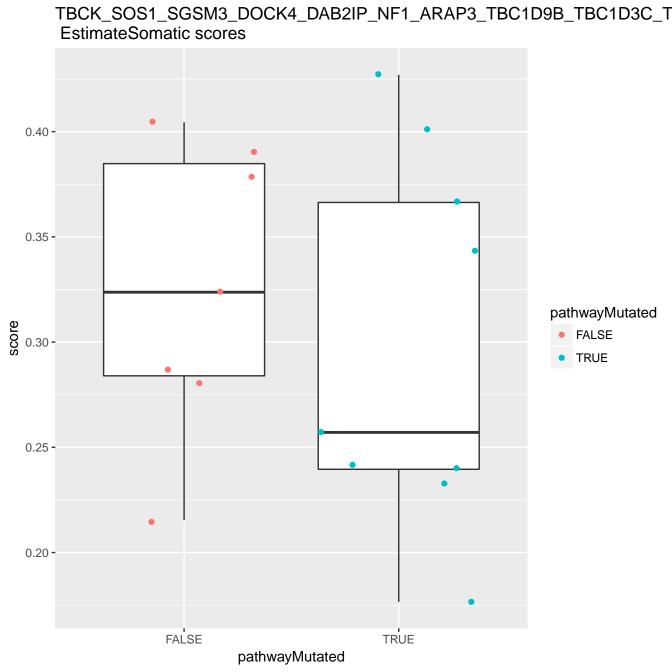


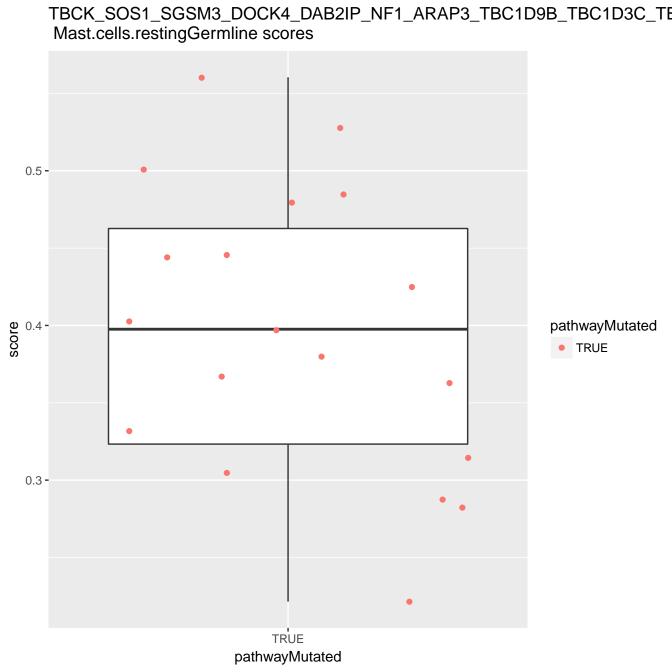


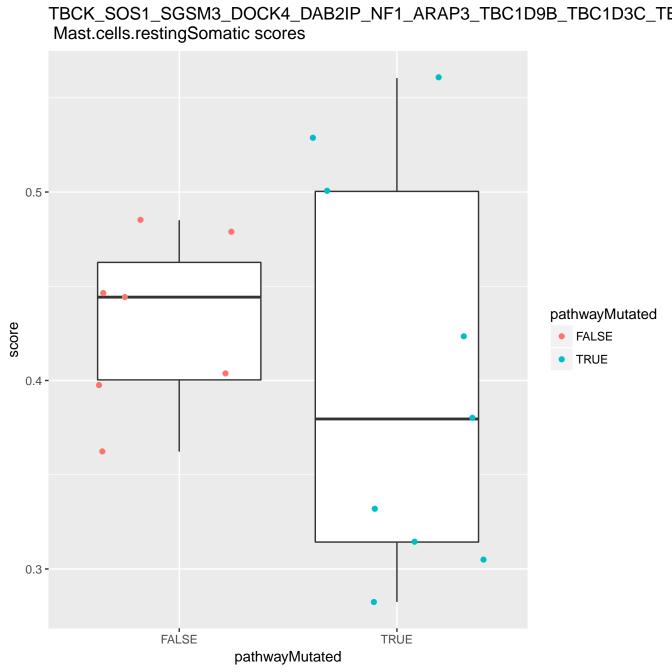


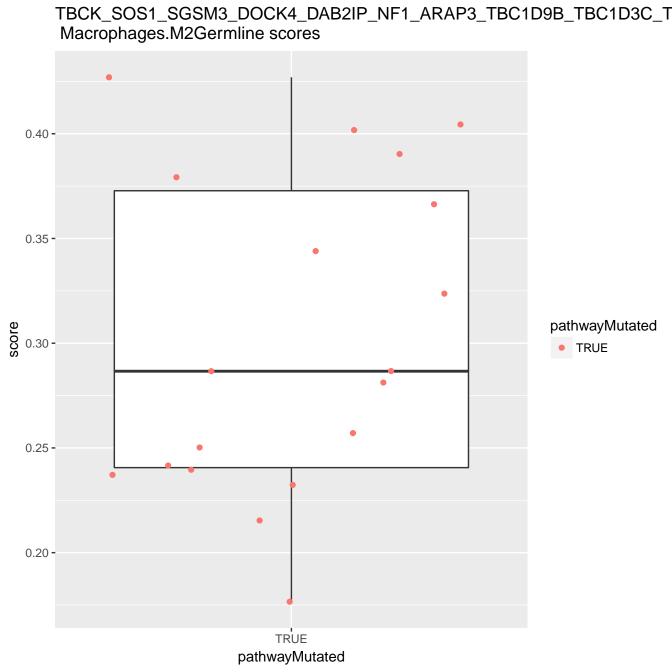
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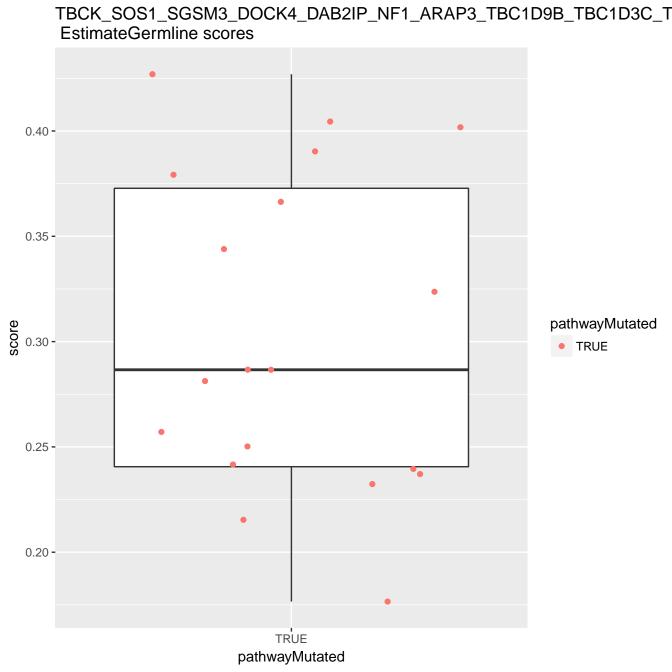


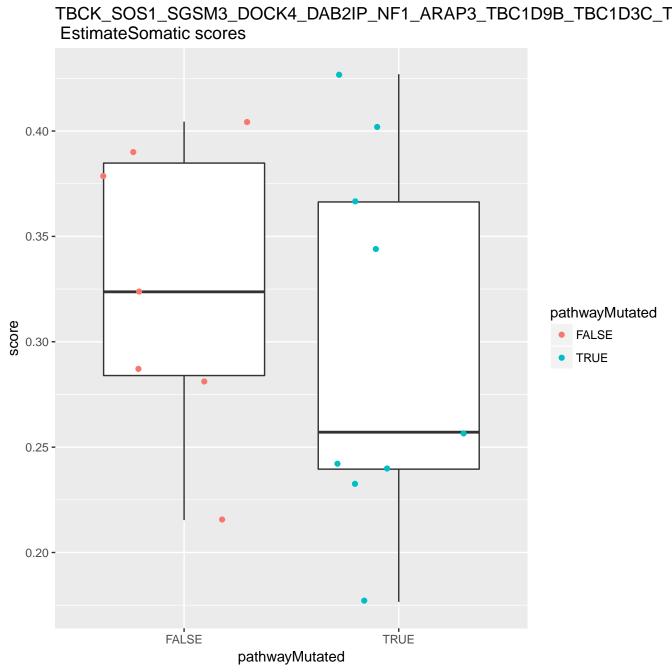


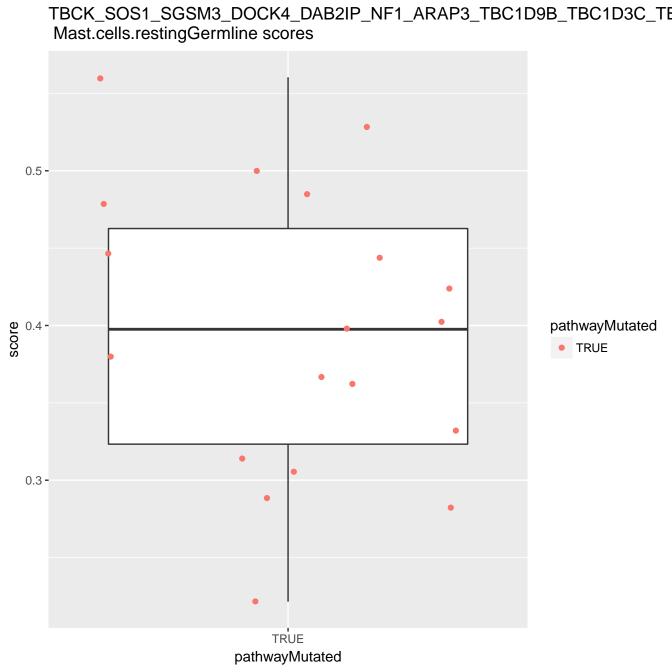


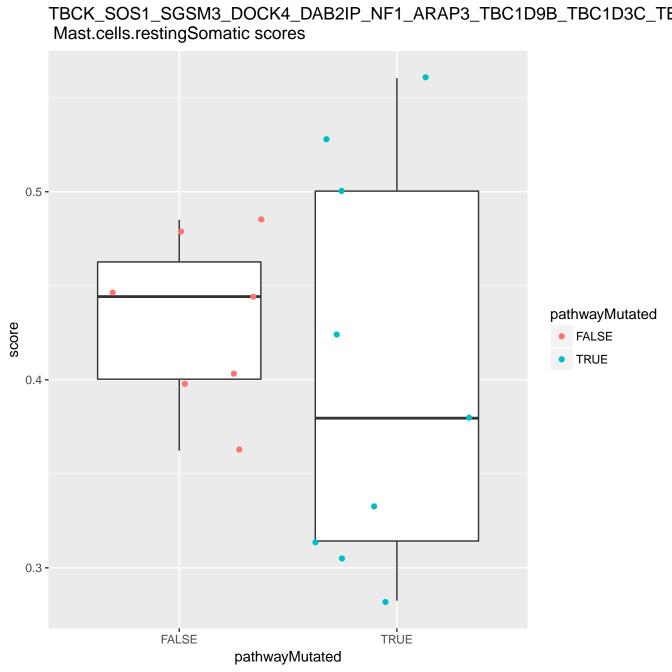


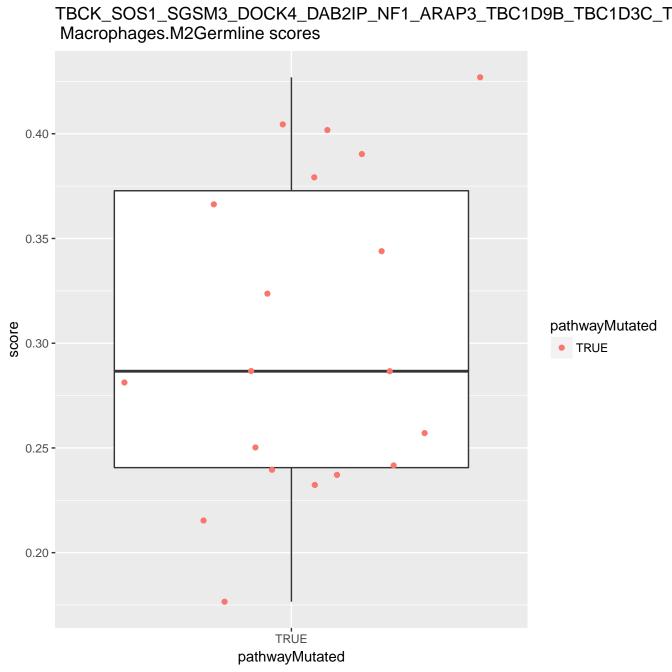
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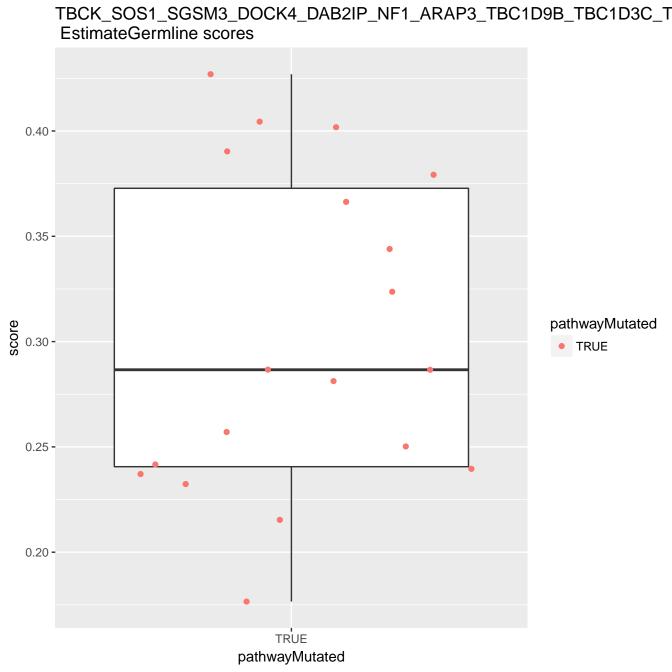


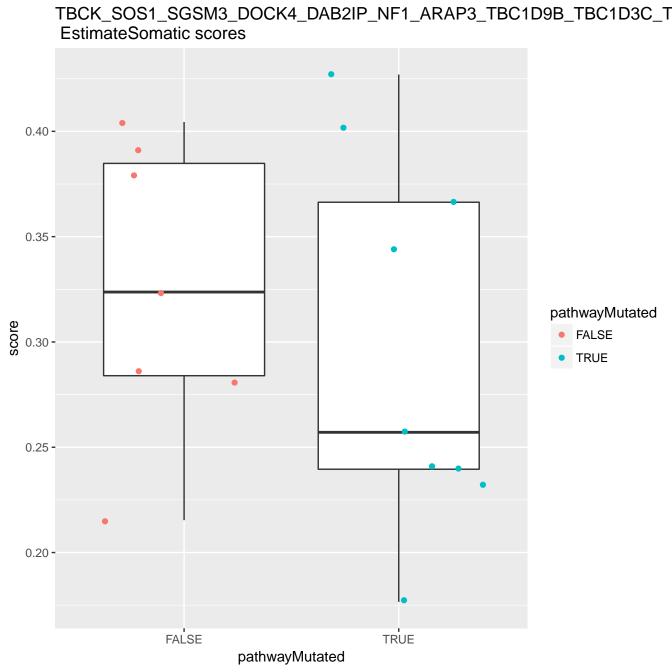


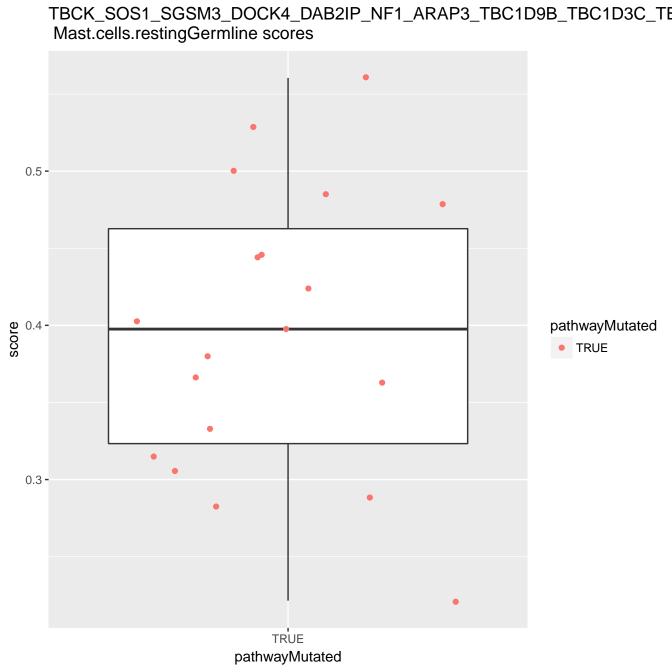


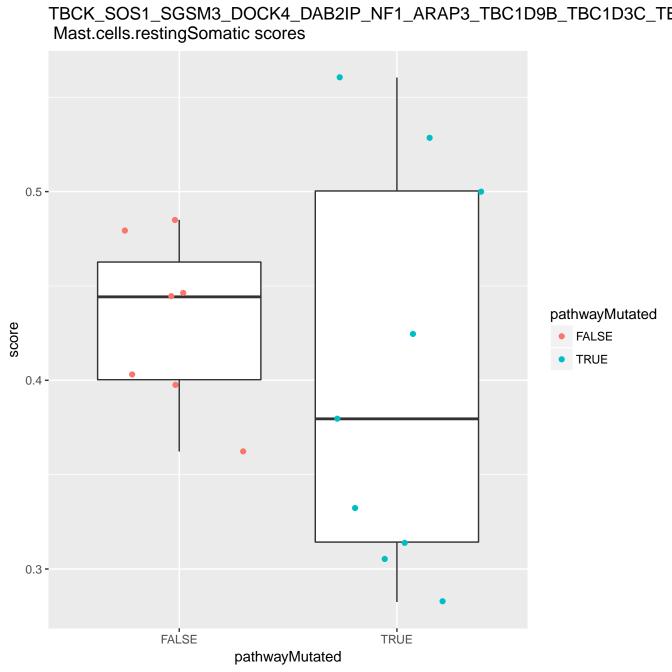


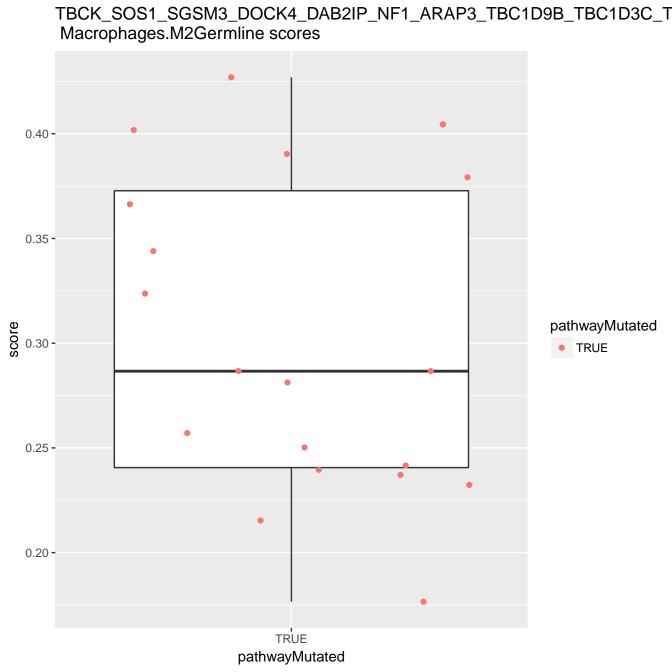
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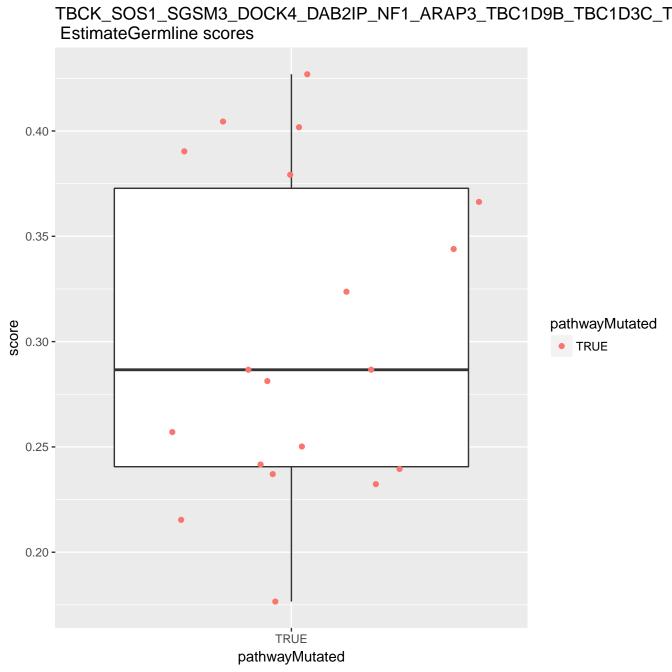


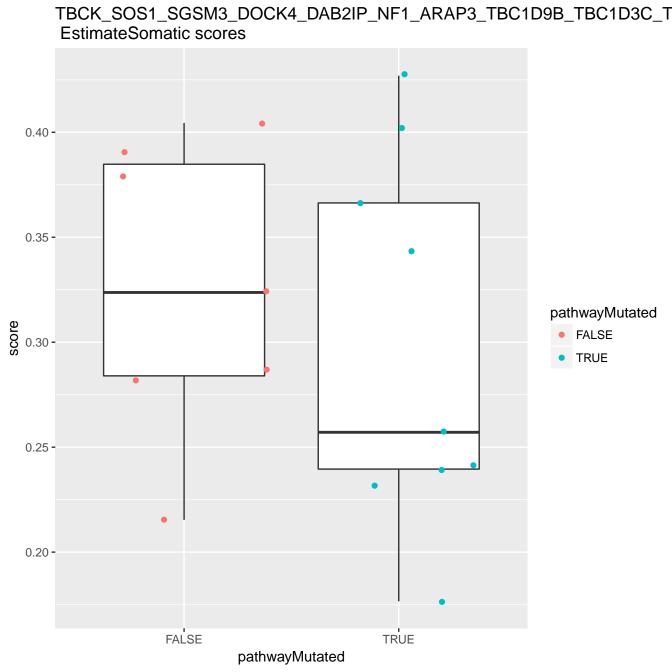


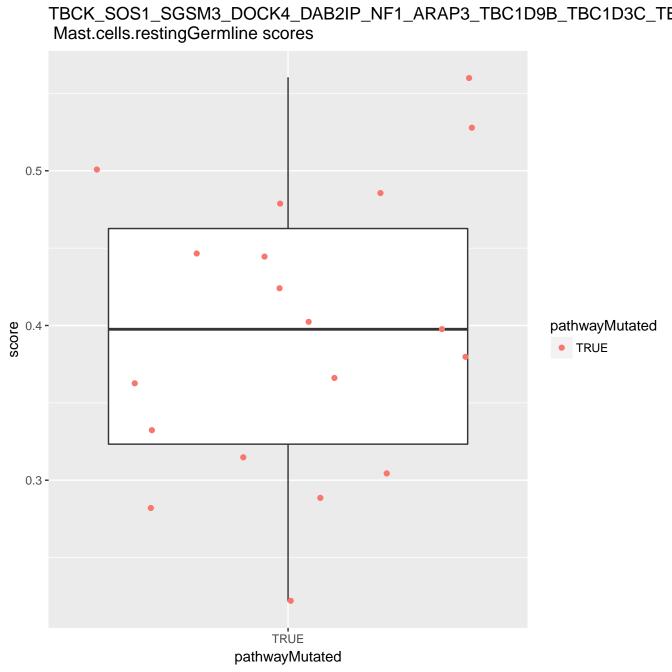


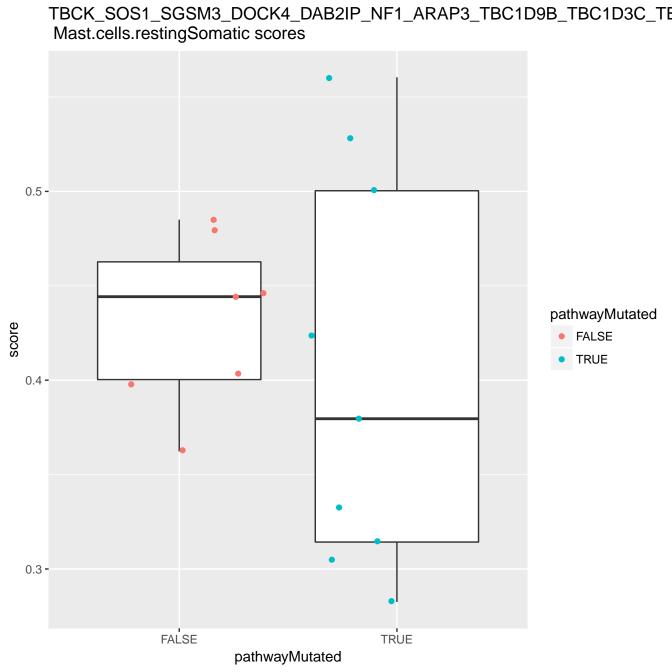


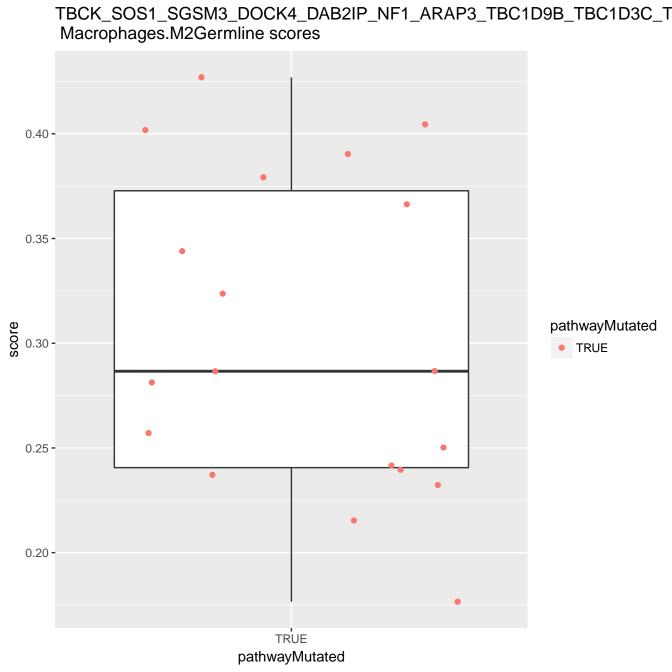
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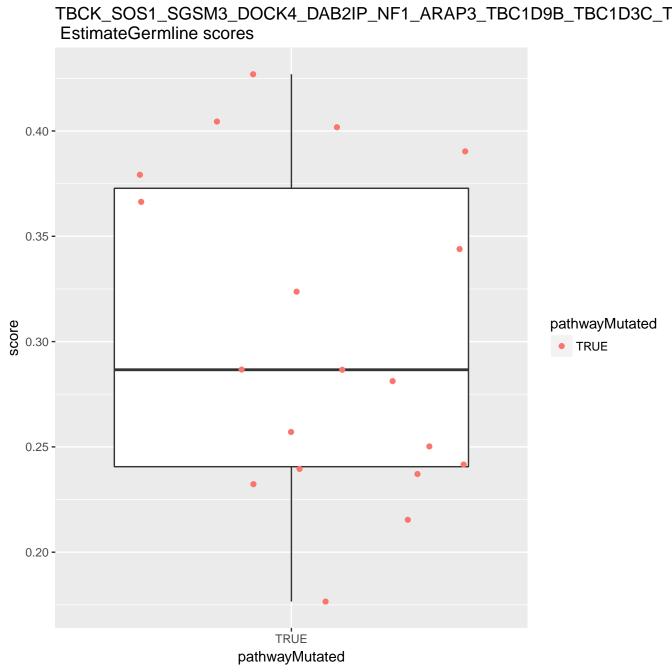


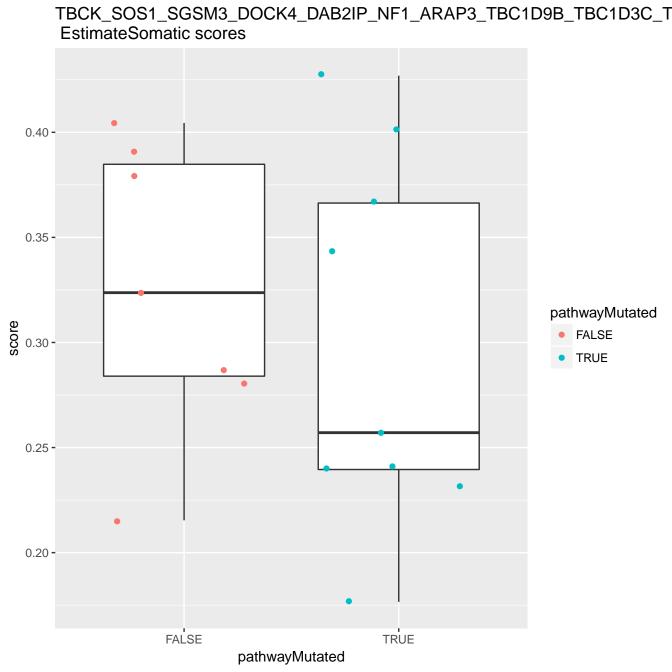


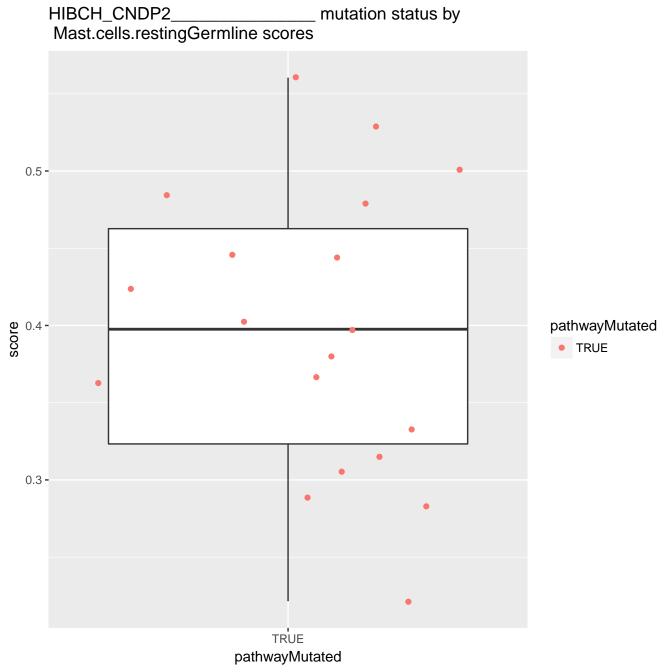


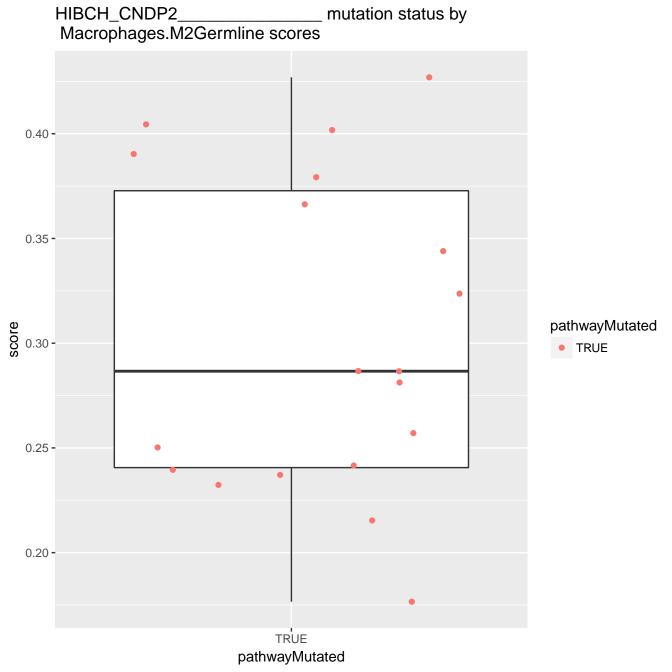


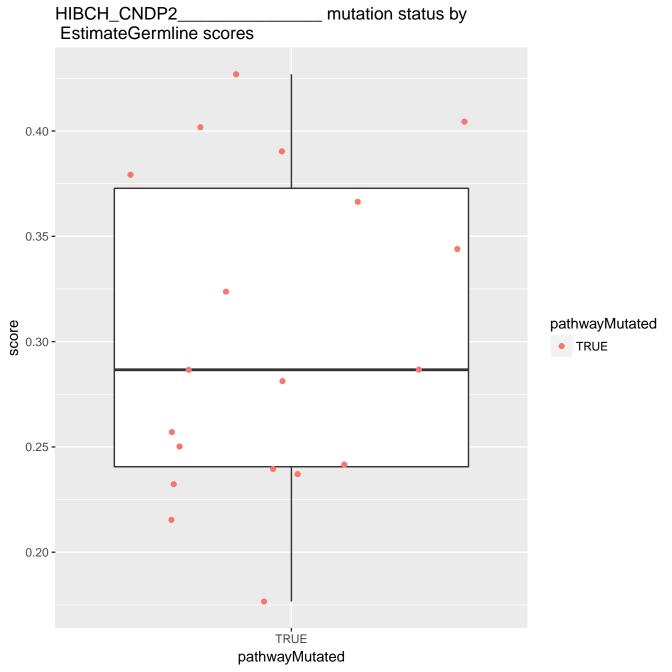
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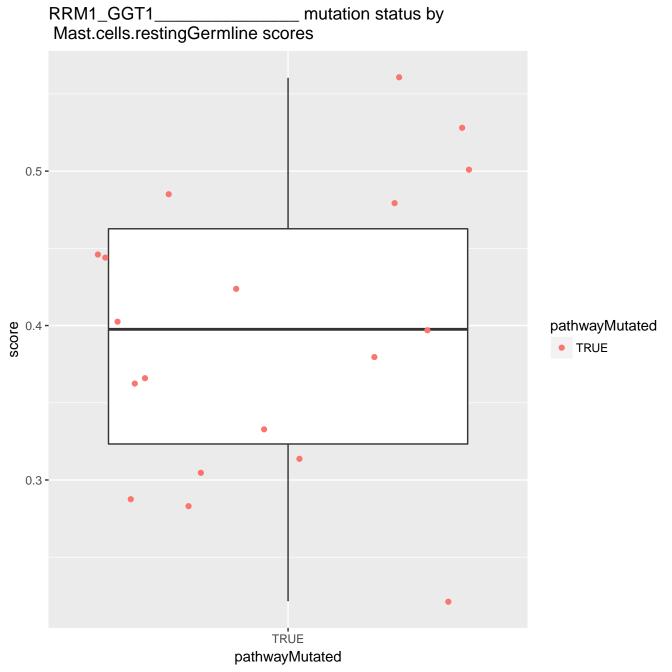


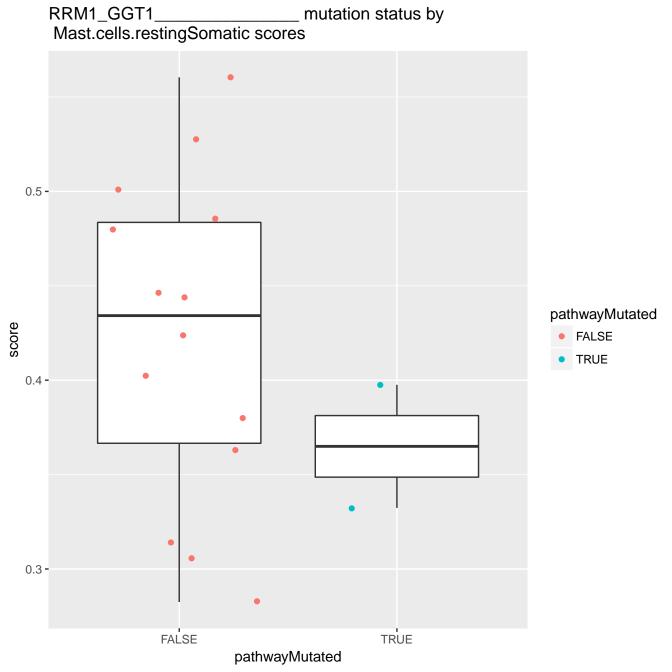


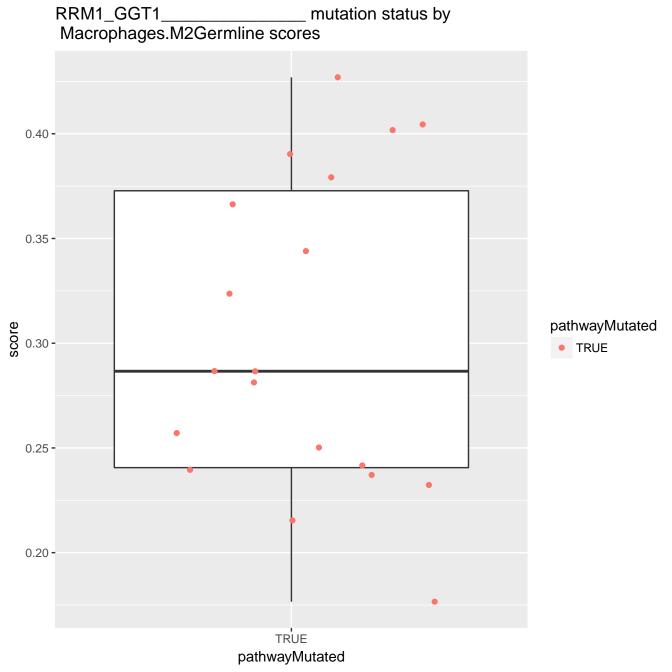


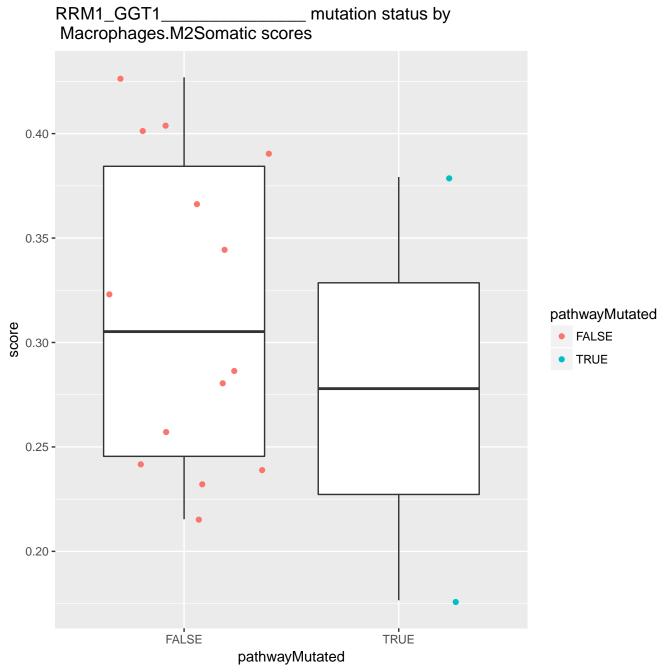


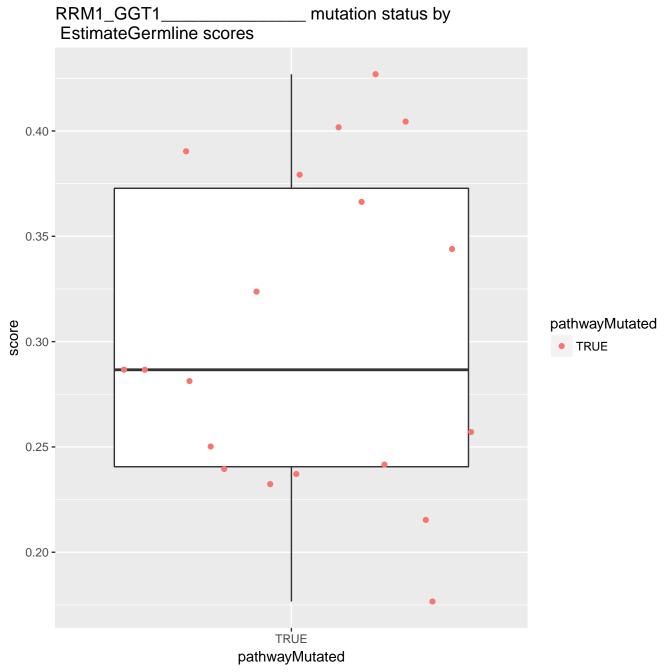


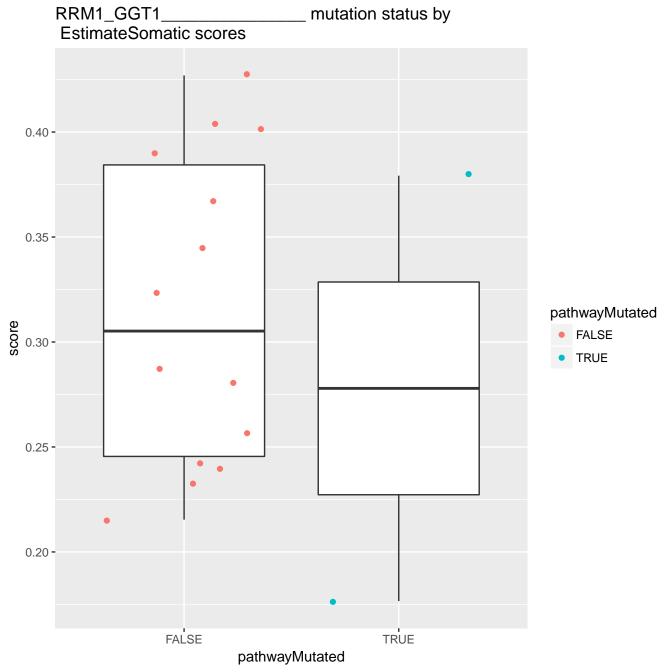


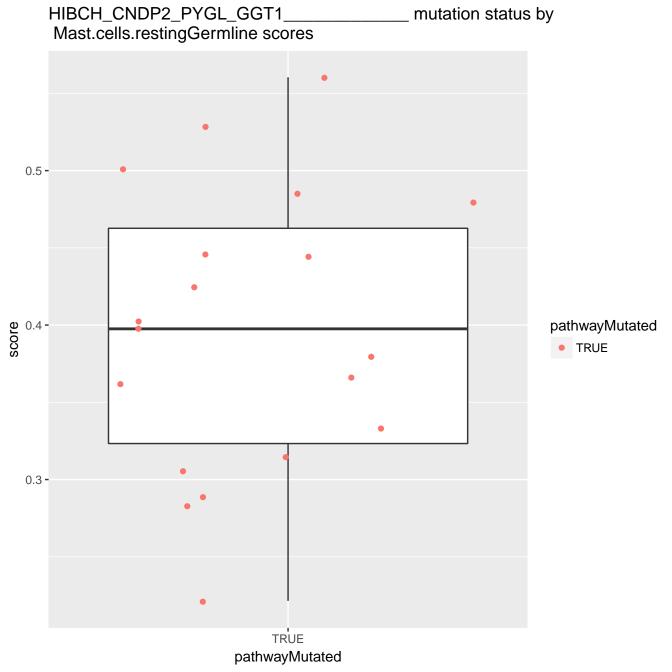


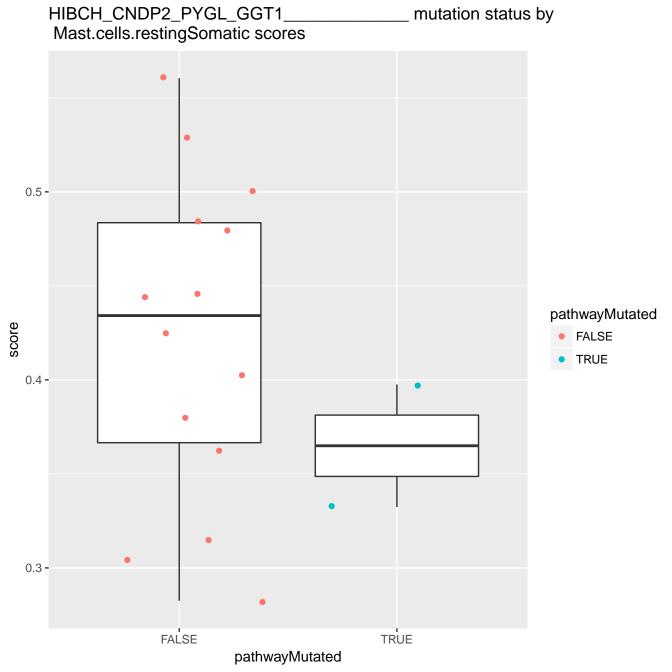


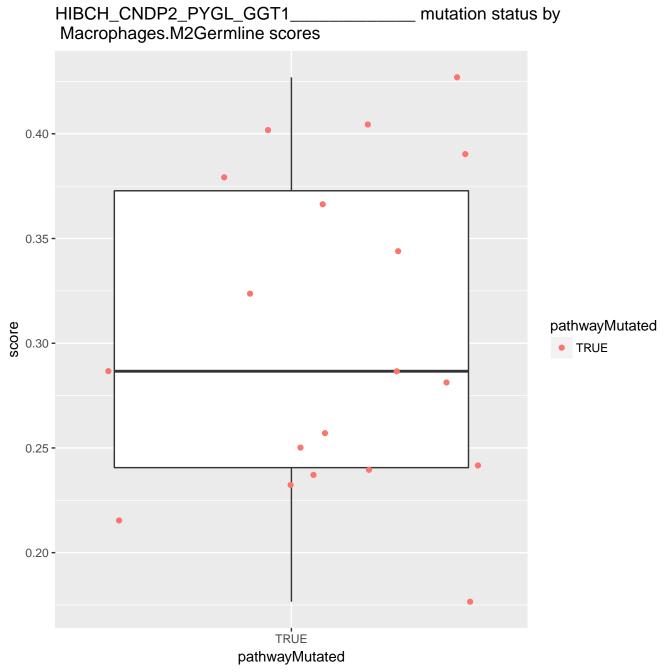


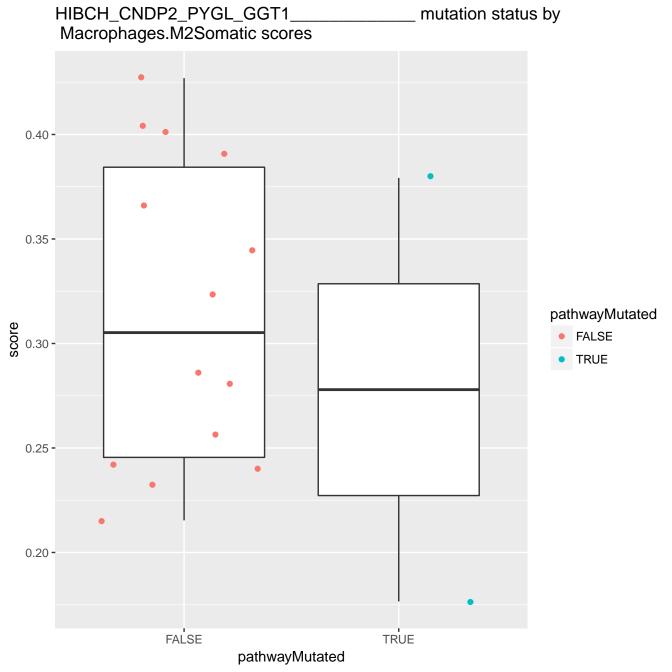


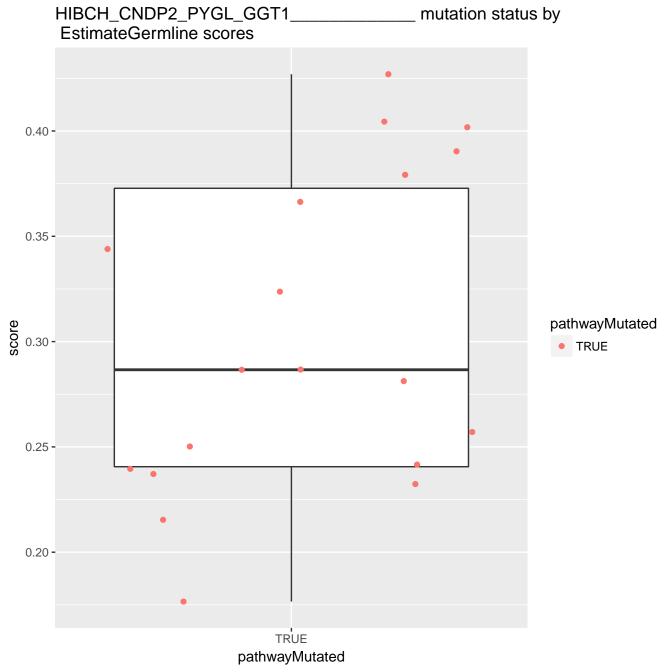


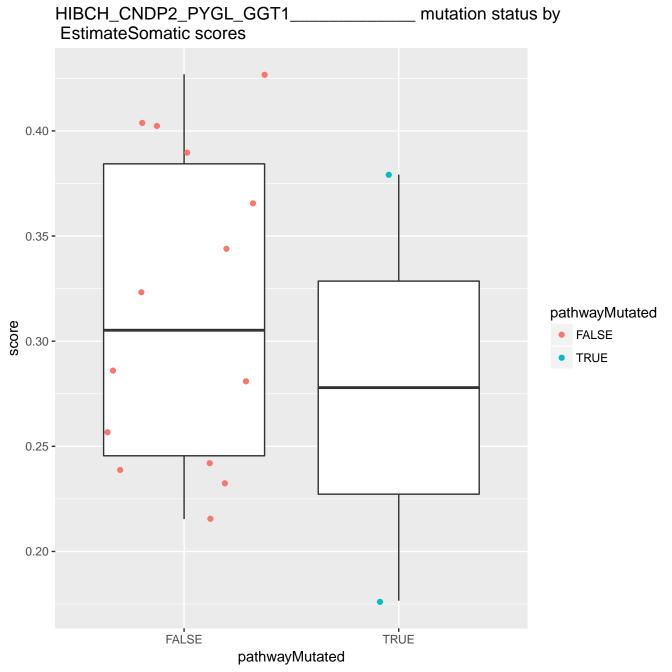


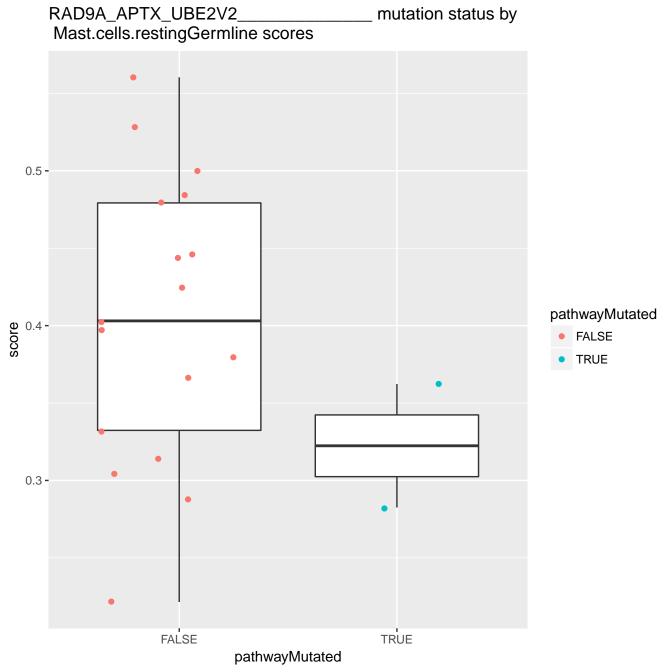


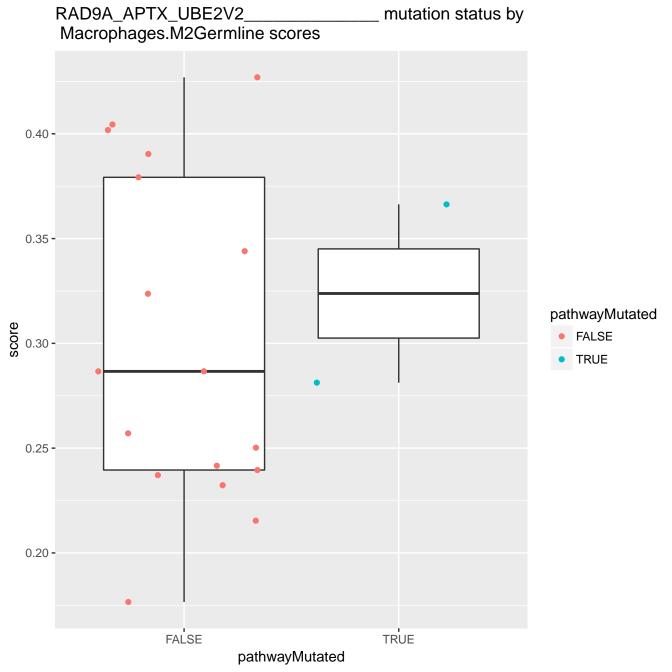


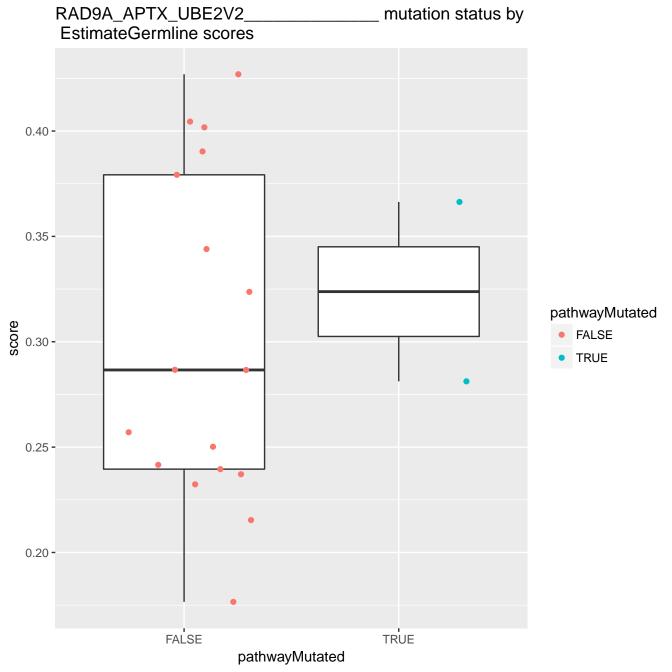


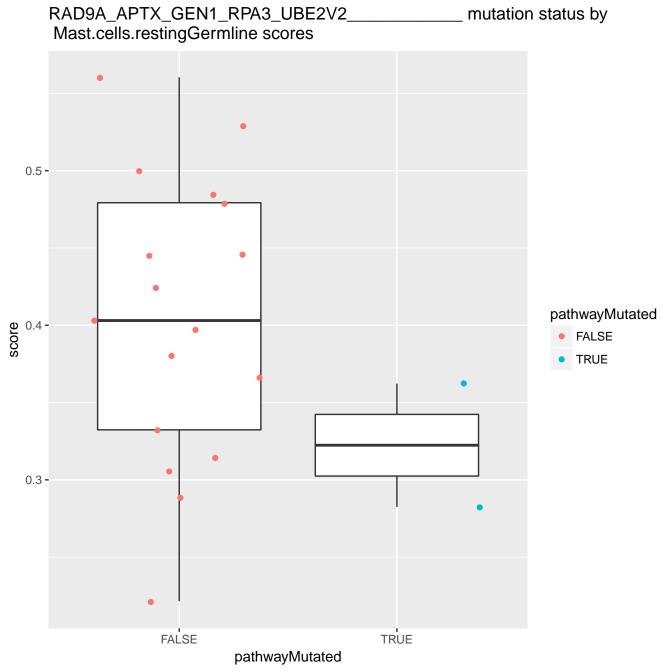


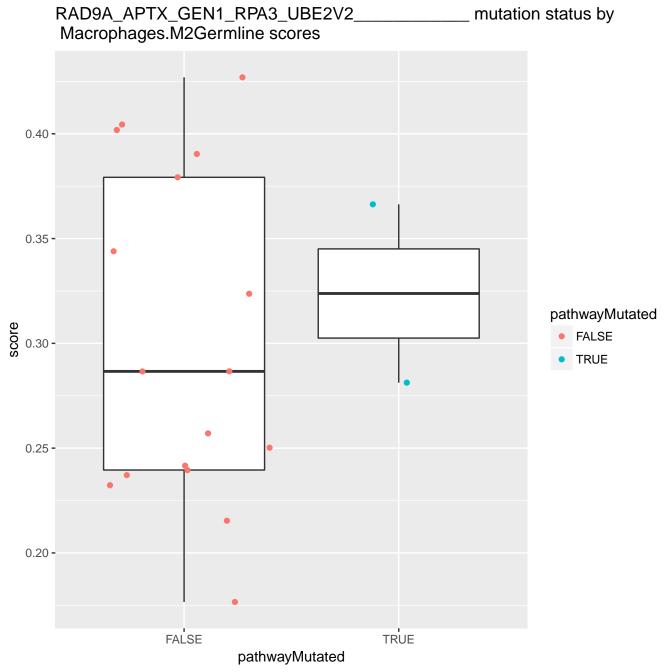


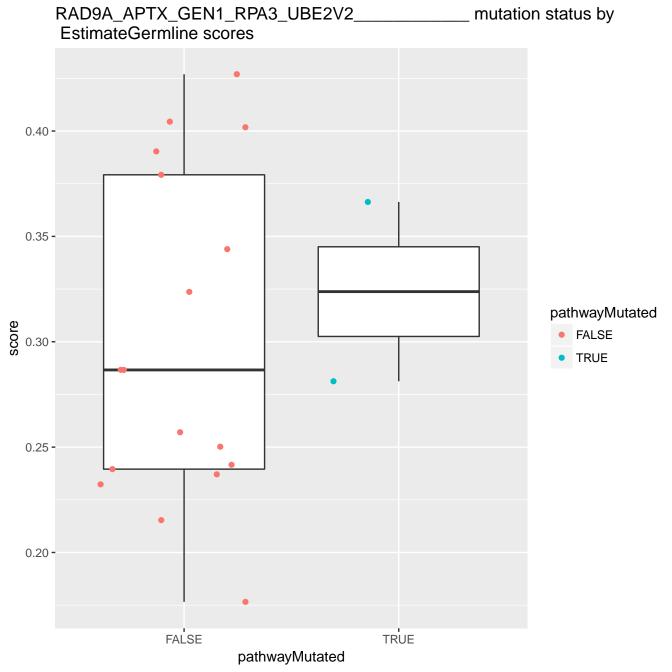












RAD9A\_APTX\_GEN1\_UBE2W\_RPA3\_RPS27A\_RAD52\_PMS2P5\_CDK9\_ Mast.cells.restingGermline scores 0.5 pathwayMutated 0.4 -FALSE TRUE 0.3 -FALSE TRUE pathwayMutated

