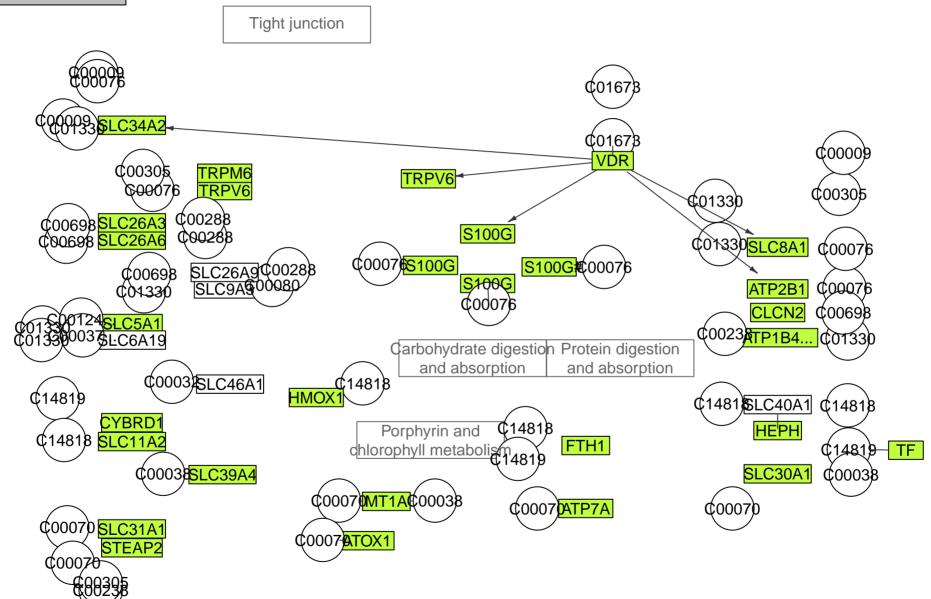
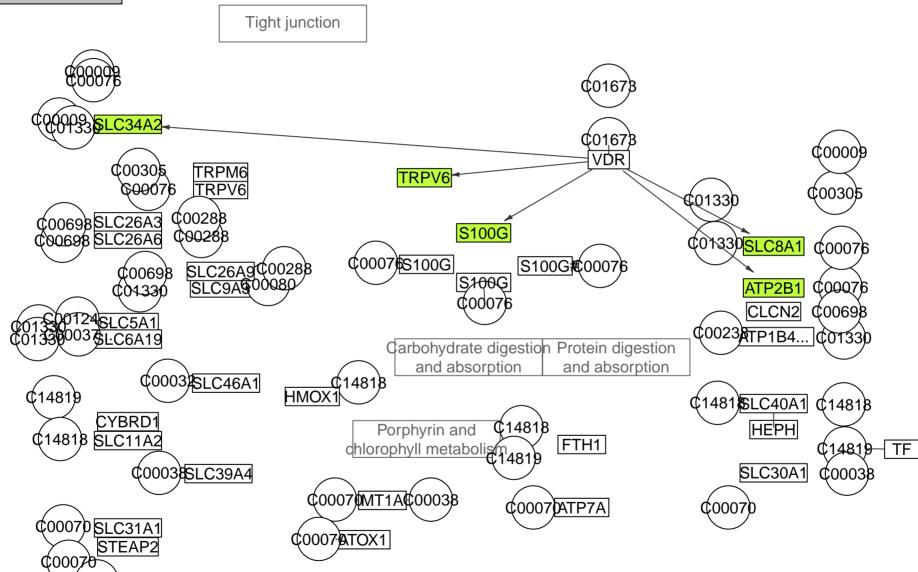
Mineral absorption (genes with data)



Mineral absorption (sink nodes)

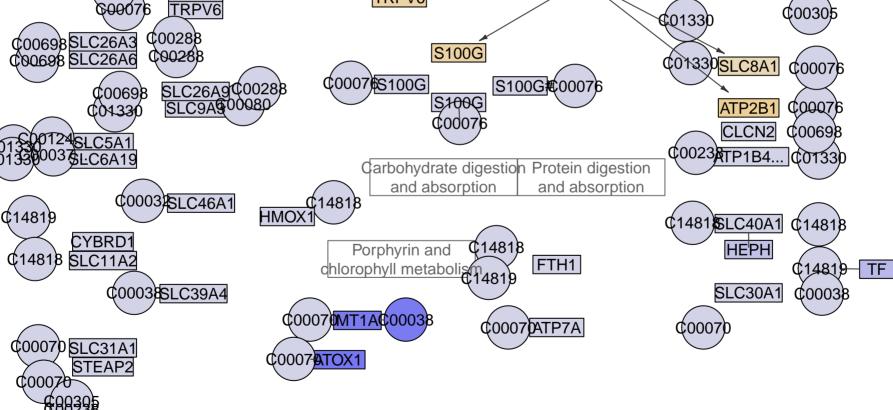


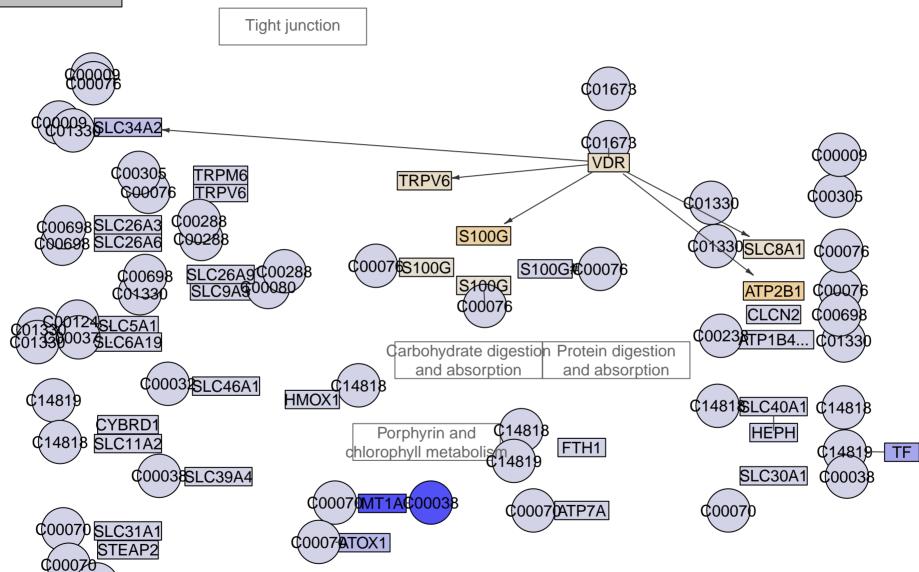
Tight junction **¢**0167 000033 **SLC34A2** C0167 ¢0000 VDR C0030 TRPM6 TRPV6 TRPV6 600076 ¢00305 ¢01330 ¢00288 C00698 SLC26A3 C00698 SLC26A6 S100G ¢90288 ¢01330SLC8A1 ¢00076 ¢0007\$\$100G SLC26A9 (0028) SLC9A (00080 S100G#00076 ¢00698 S100G ATP2B1 C00076 **¢**04330 C00076 CLCN2 C00698 00124SLC5A1 ¢0023 003 \$LC6A19 C01330 Carbohydrate digestion Protein digestion and absorption and absorption C0003 SLC46A1 ¢14819 HMOXC1481 SLC40A1 C14818 CYBRD<sup>2</sup> C14818 Porphyrin and HE<sub>PH</sub> C14818 SLC11A2 chlorophyll metabolig (1481) FTH1 ¢14819 TF C0003 SLC39A4 SLC30A1 ¢00038 C00070MT1AC00038 C0007)ATP7A (0007) ¢00070 SLC31A1 **C**0007**ATOX1** STEAP2 ¢0007<del>0</del>

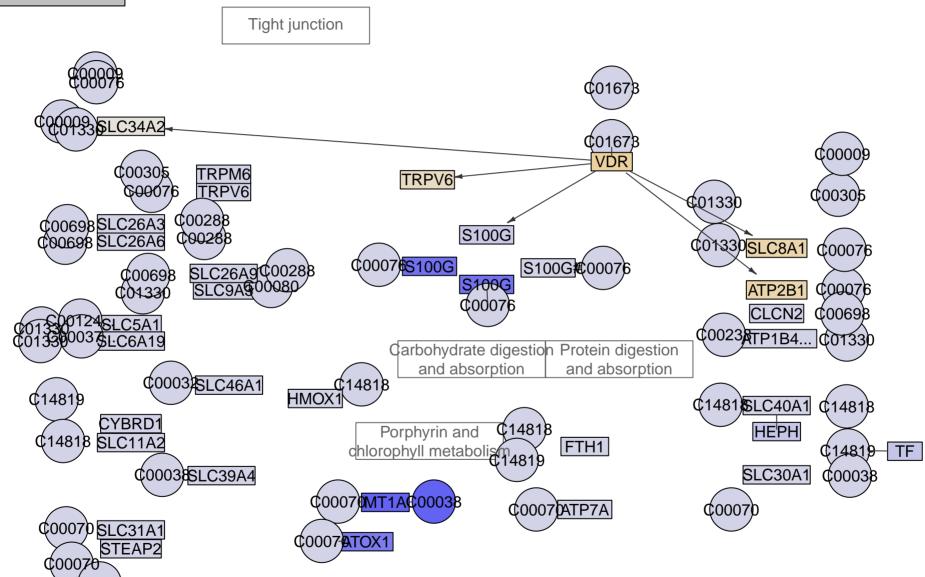
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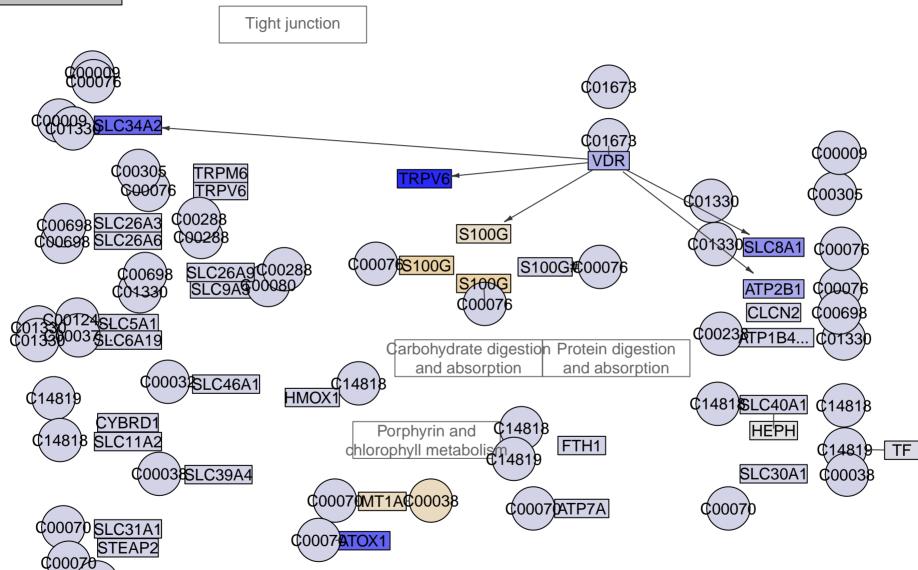
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Tight junction **¢**0167 C0167 ¢0000 VDR TRPM6 TRPV6 TRPV6 ¢00305 ¢01330 S100G ¢01330<mark>SLC8A1</mark> ¢00076 ¢0007\$\$100G S100G#00076 S100G ATP2B1 C00076 C00076 CLCN2 C00698 ¢0023 C01330 Carbohydrate digestion Protein digestion and absorption and absorption

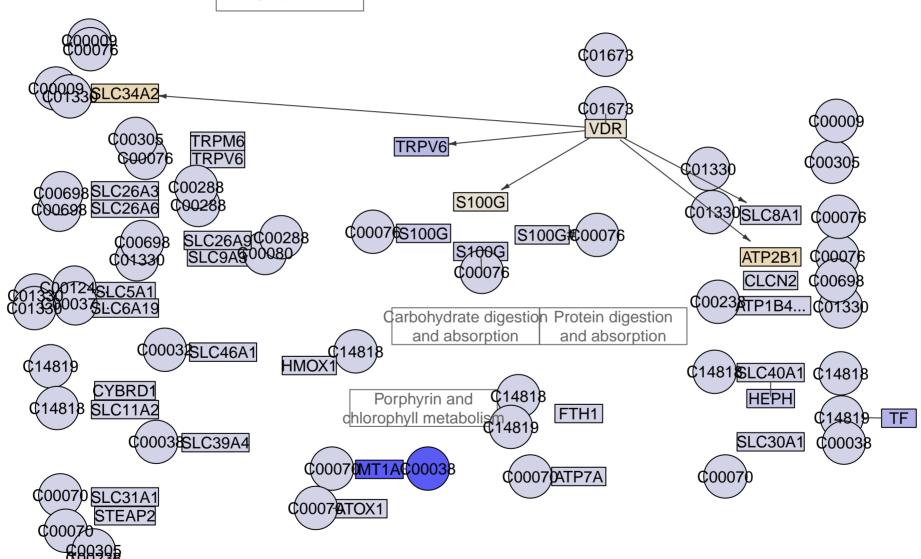


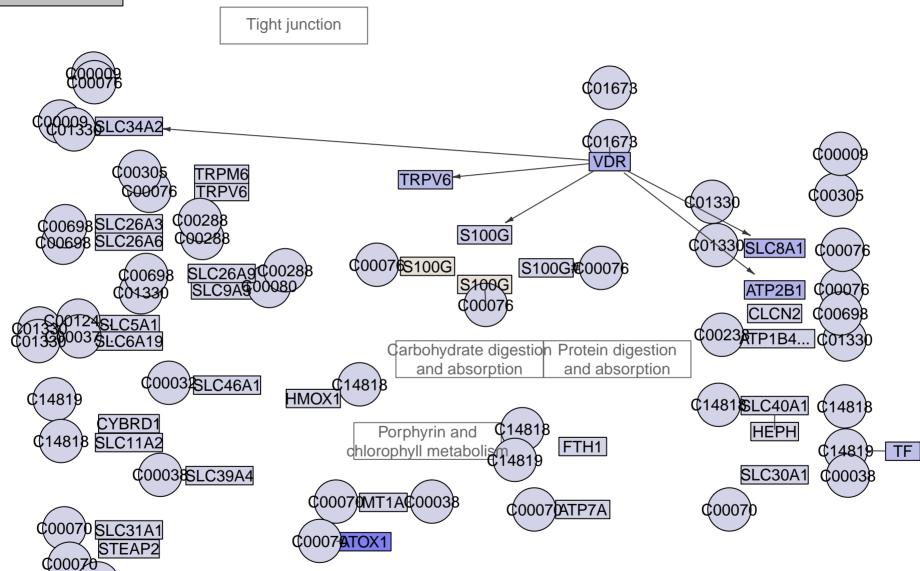


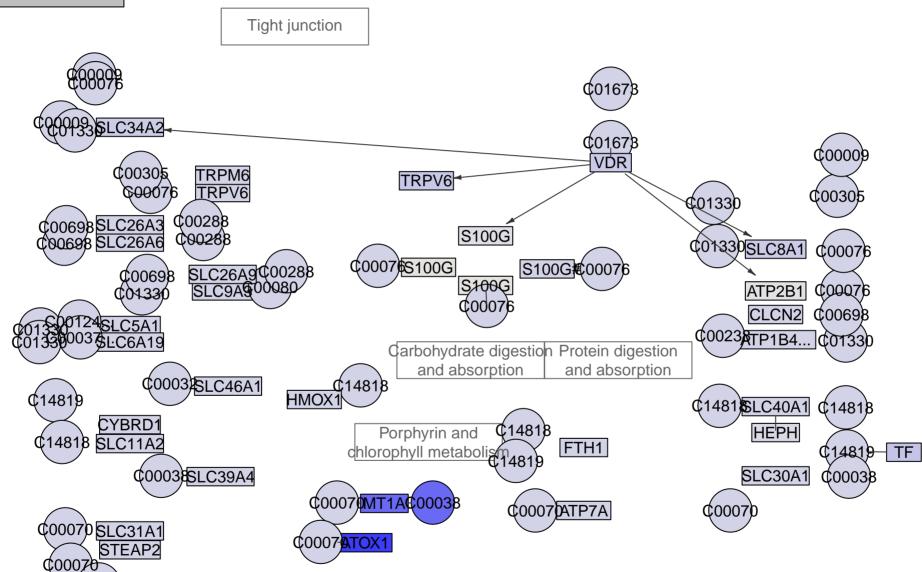




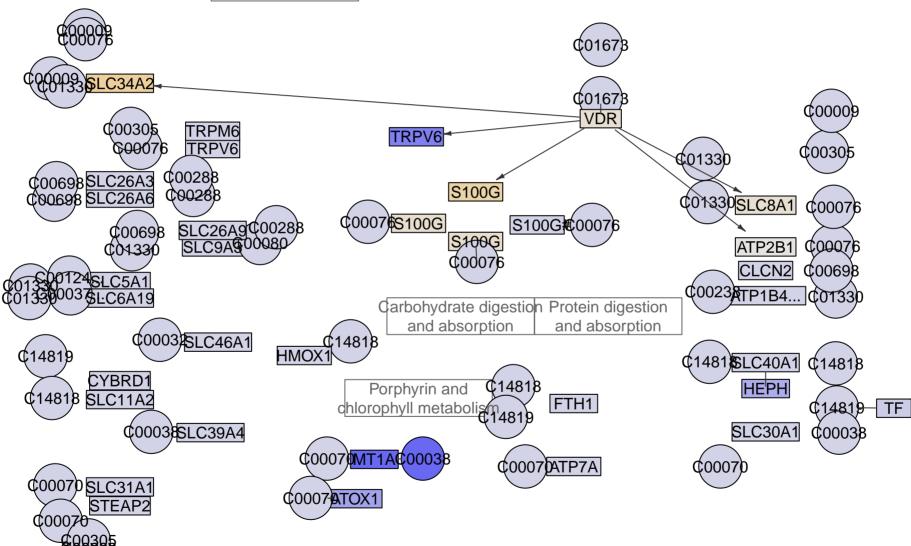
Tight junction







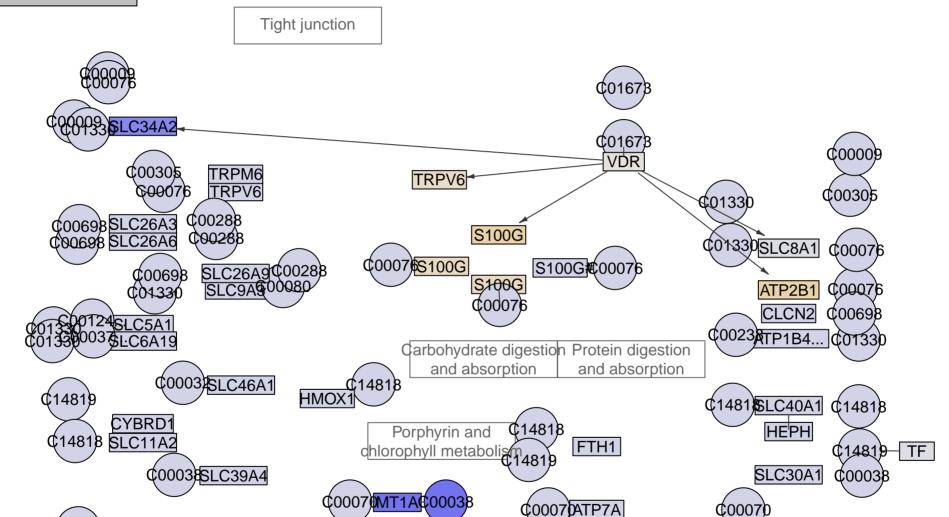




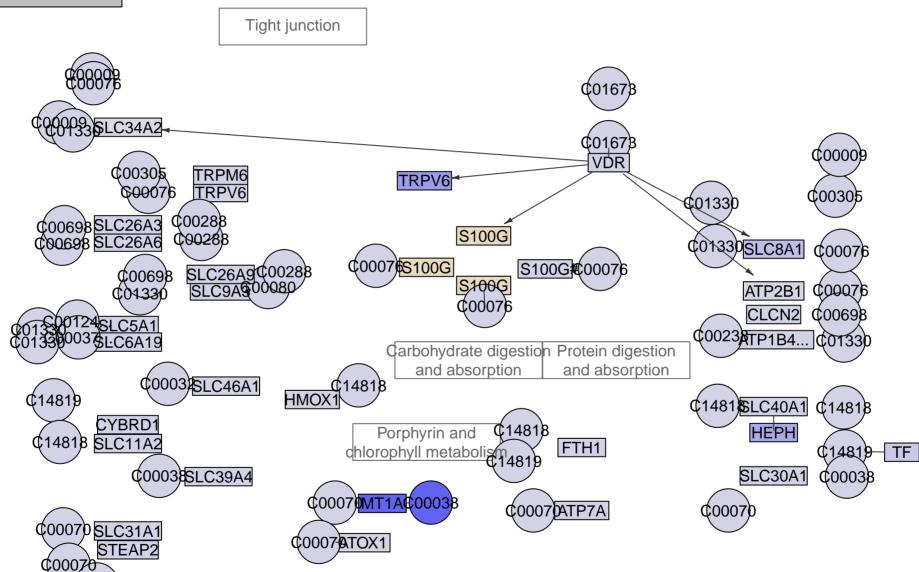
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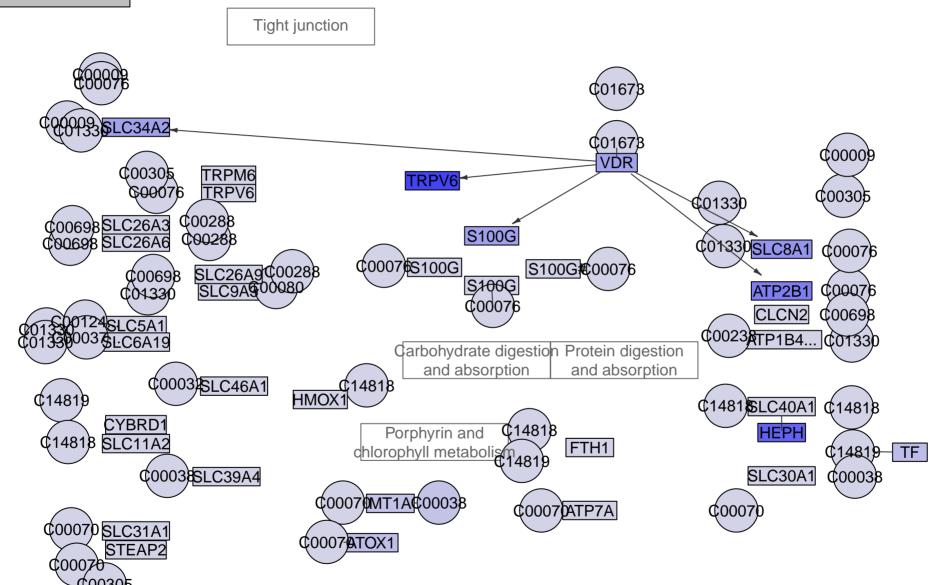
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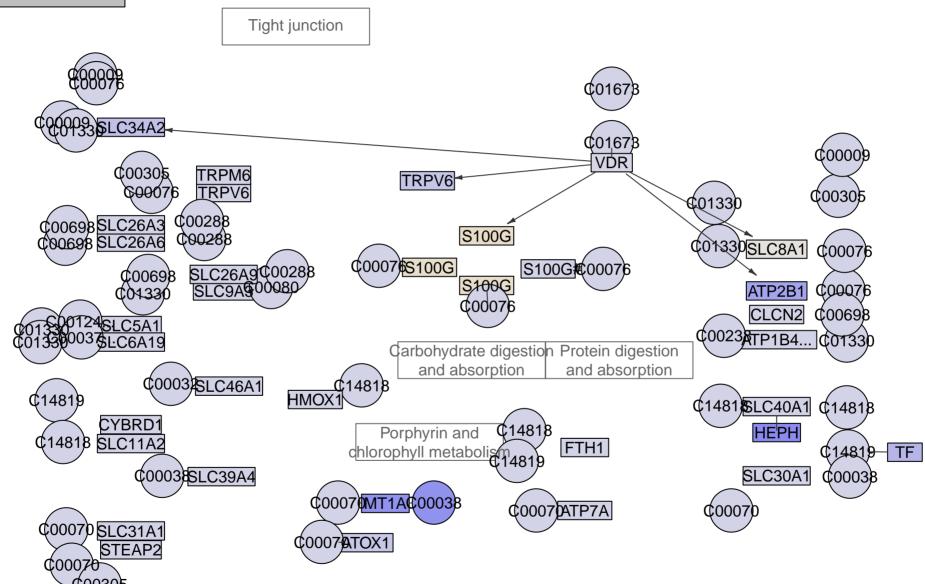
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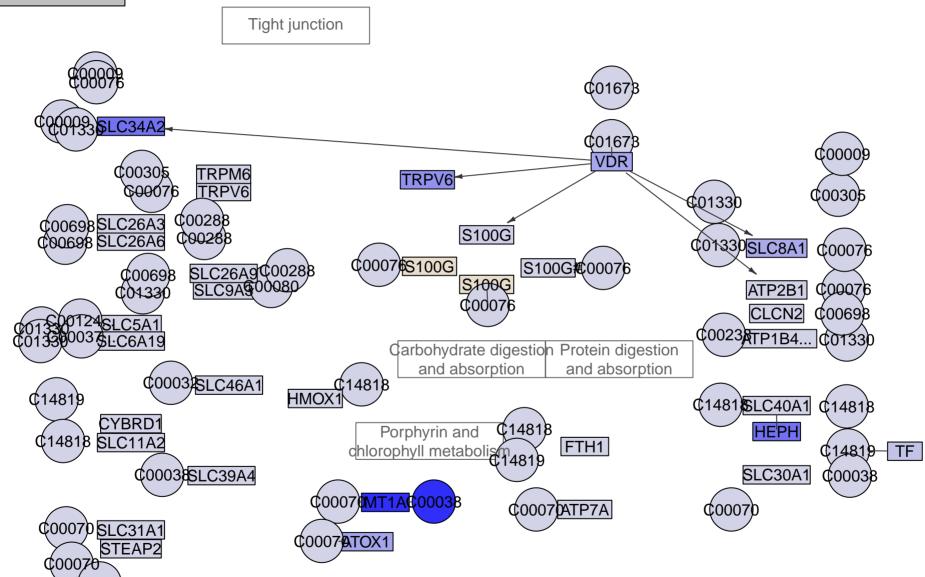


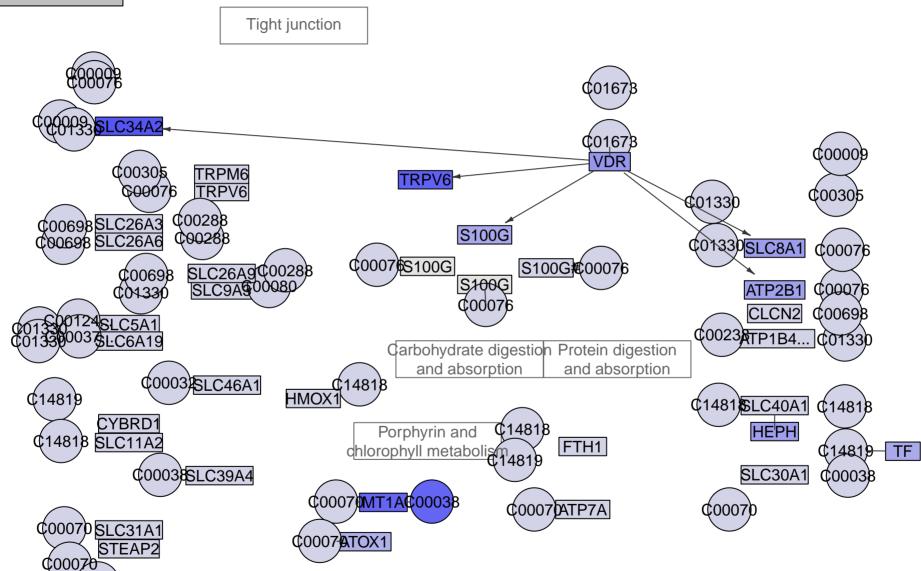
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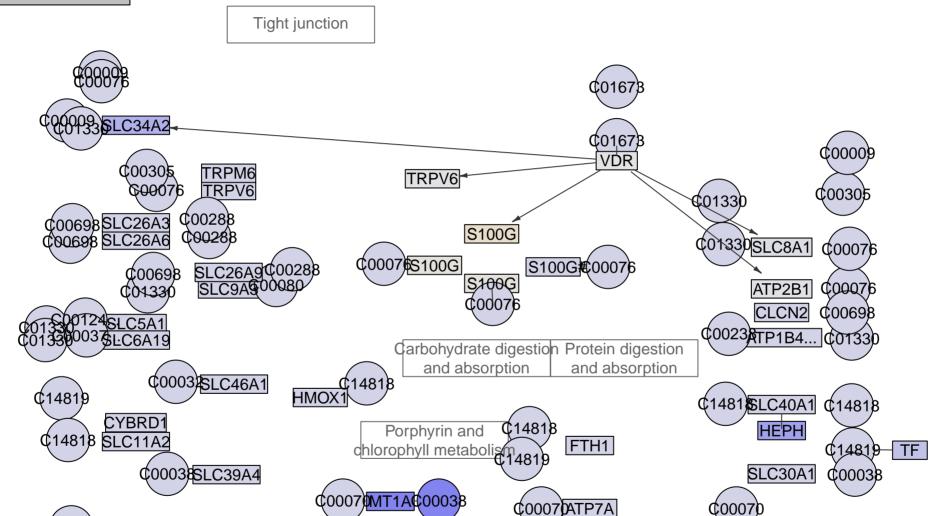




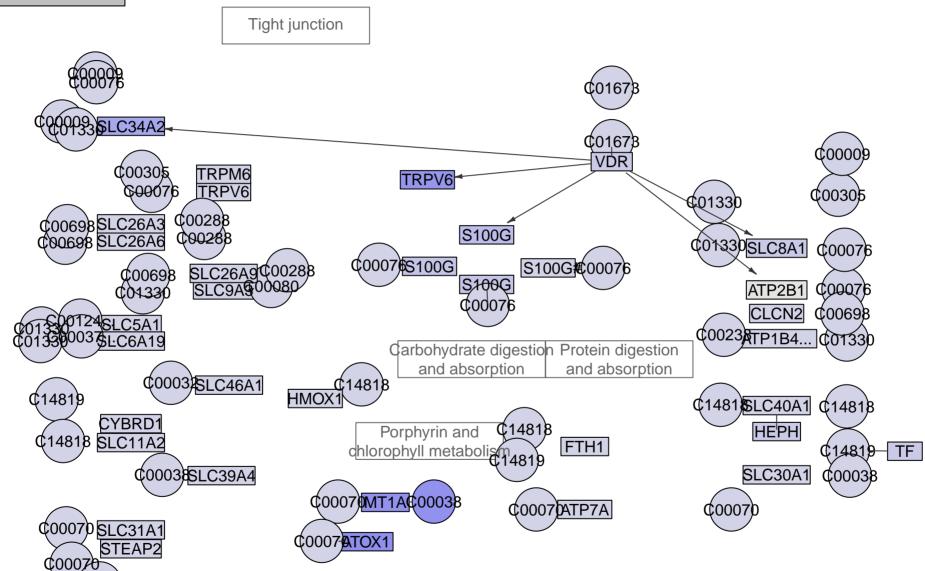
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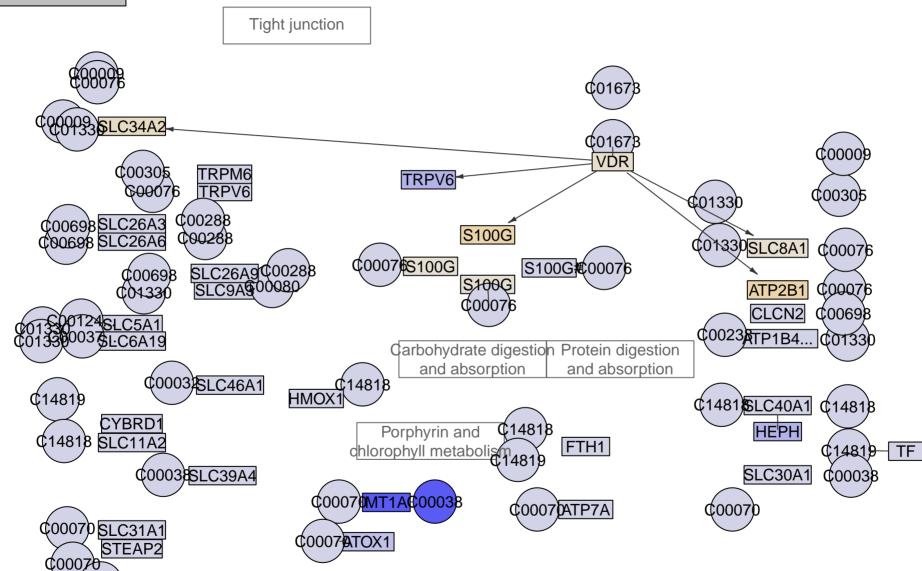
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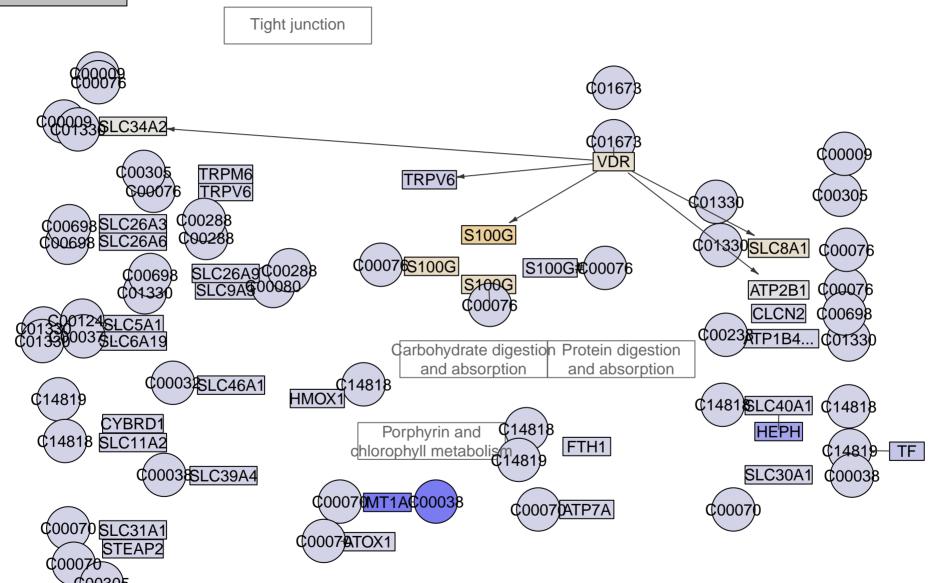
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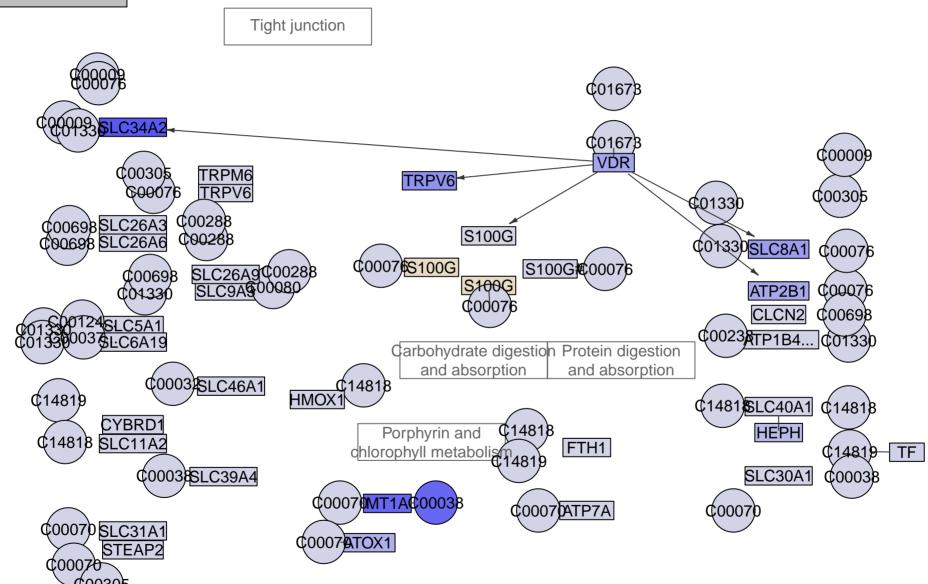


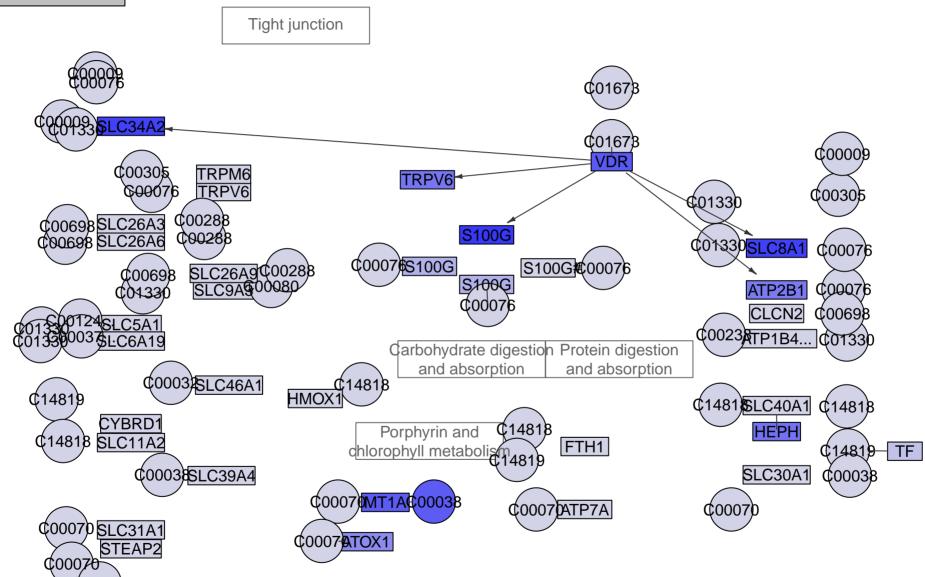
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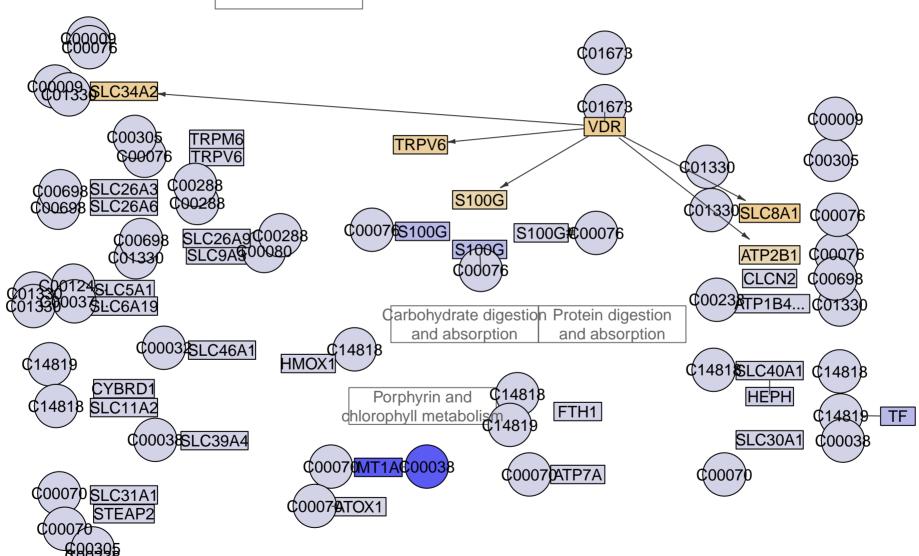


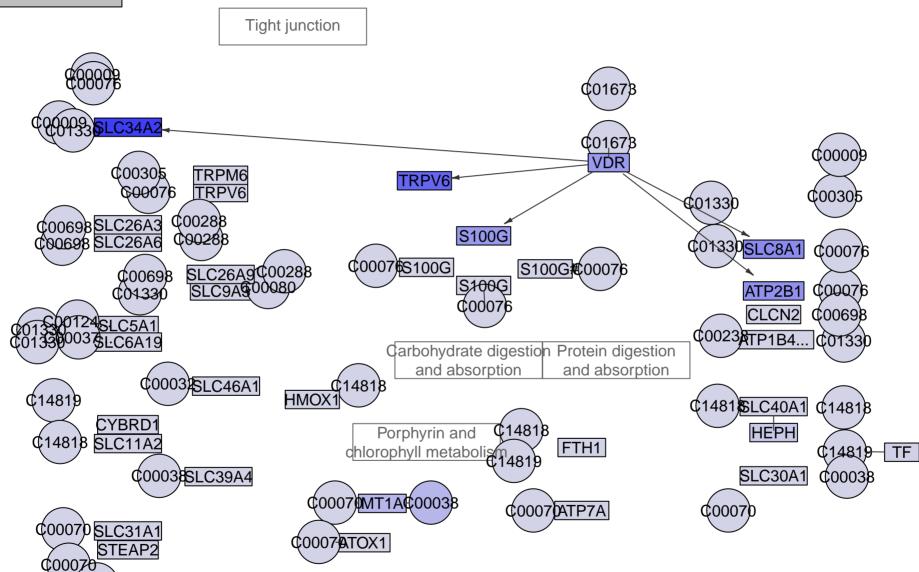


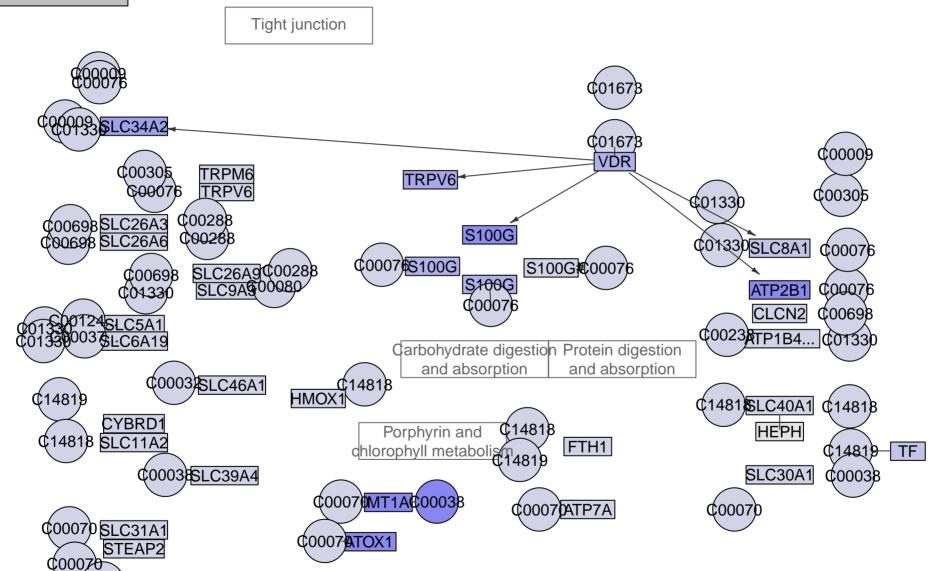


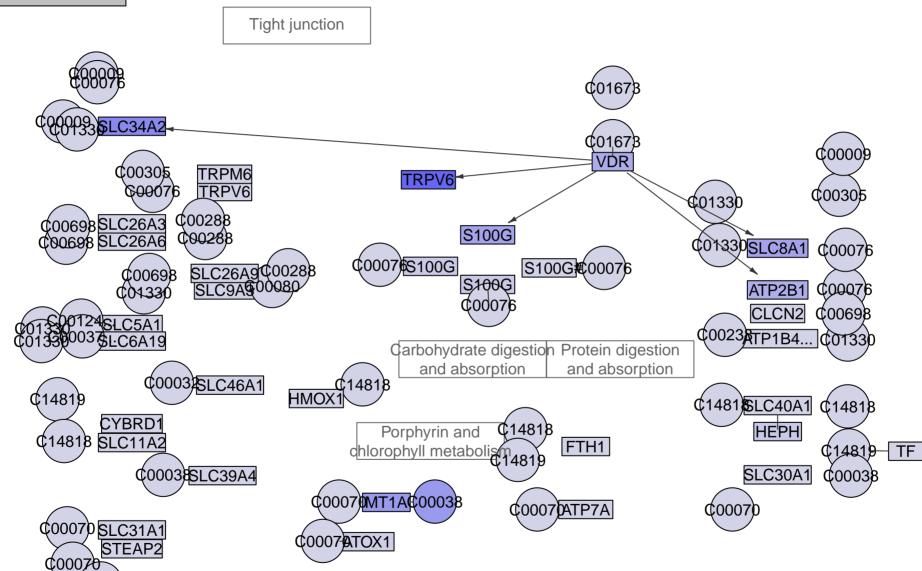


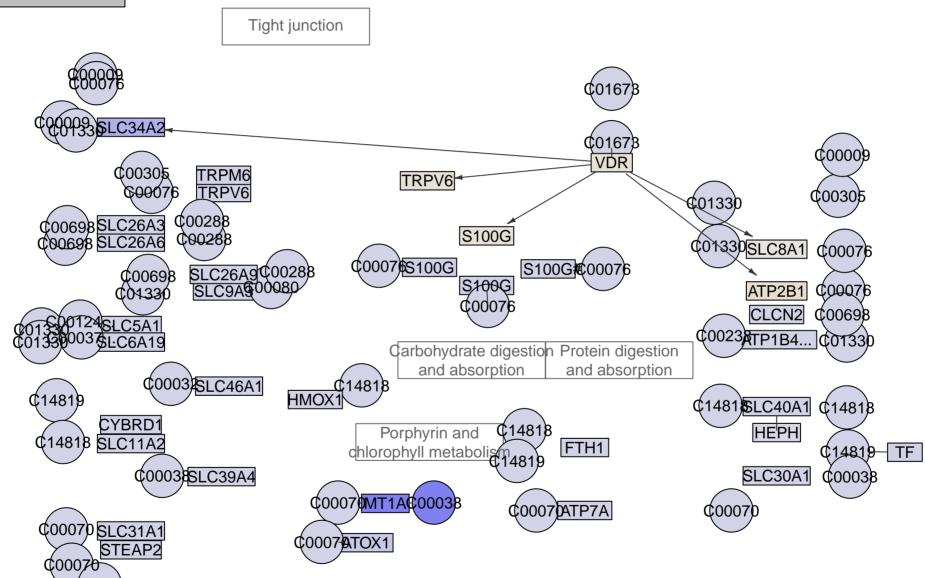




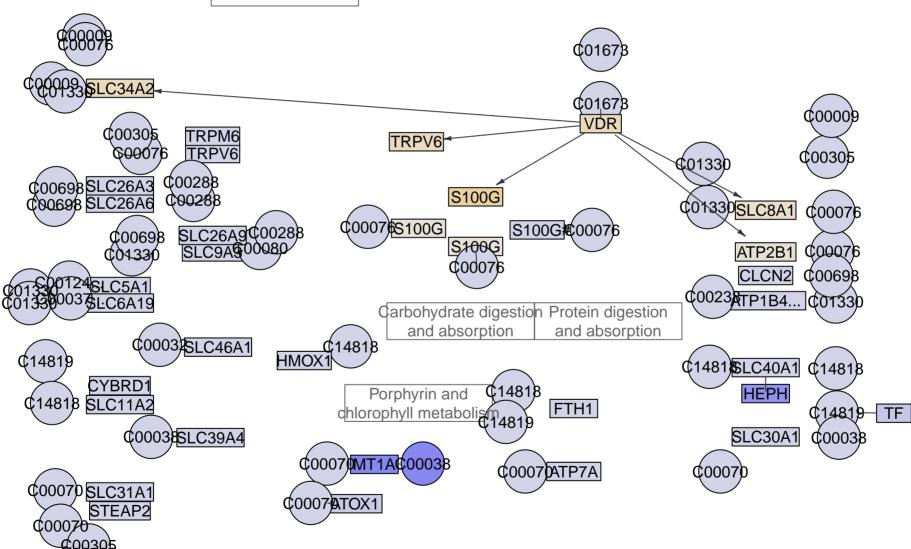


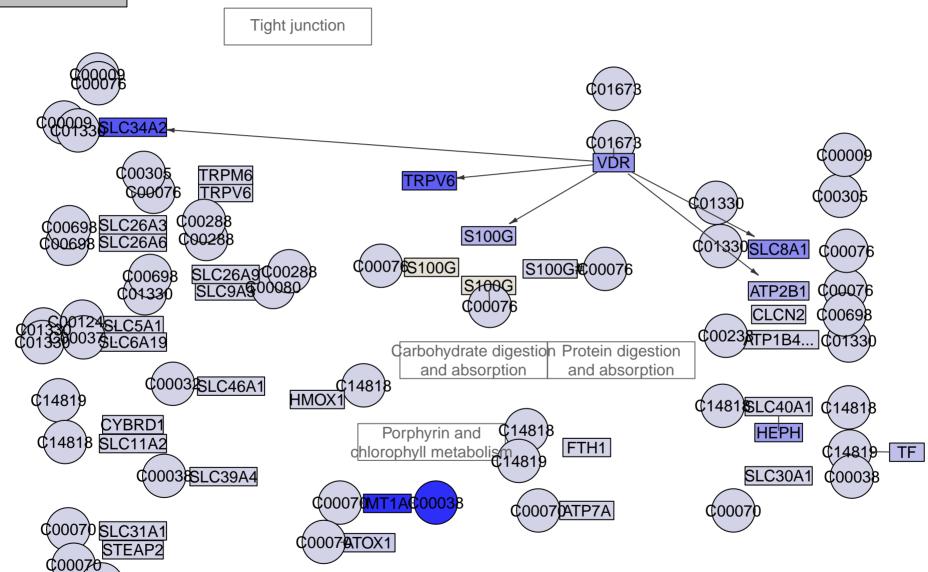






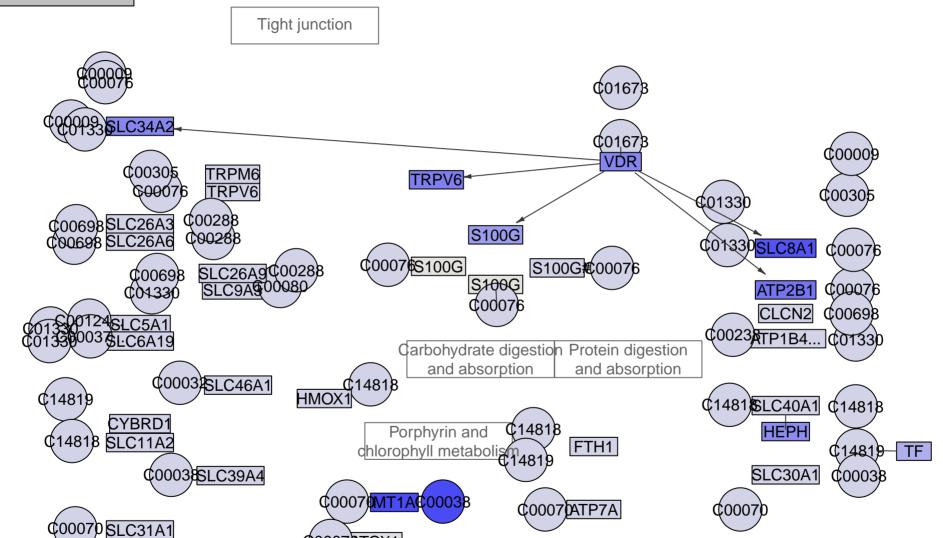




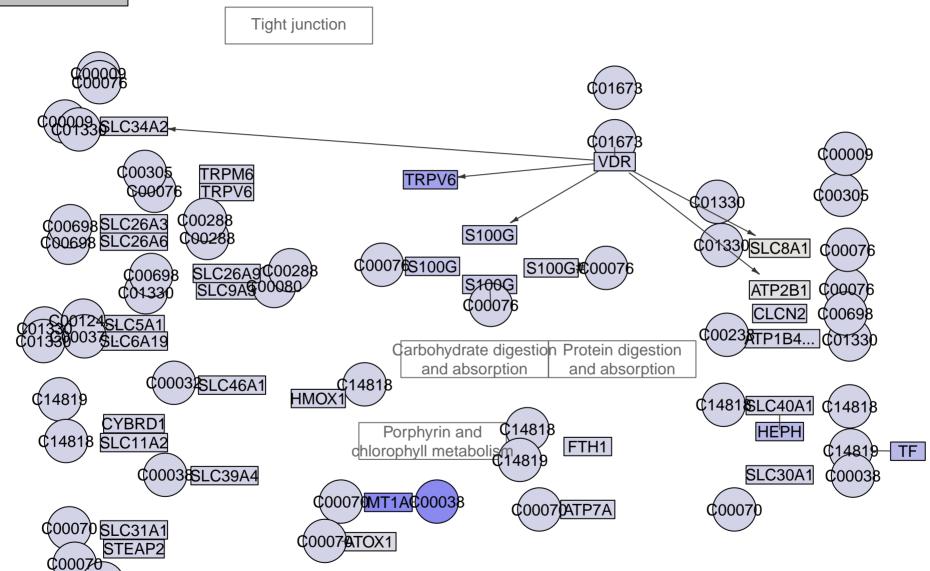


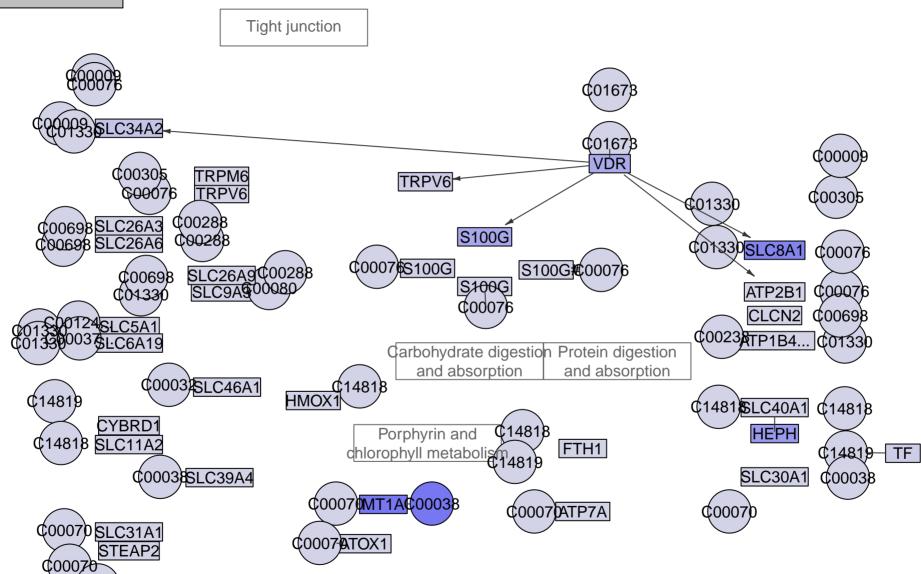
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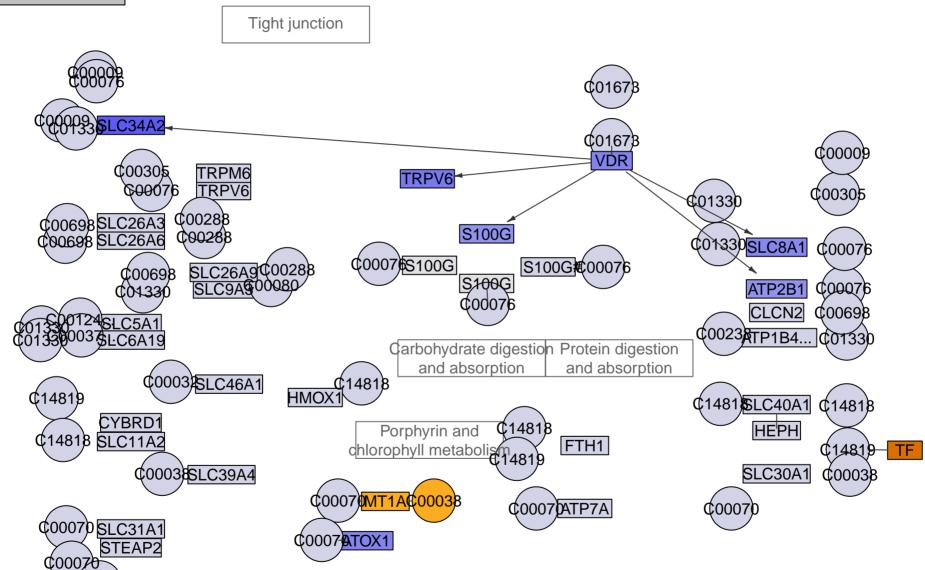
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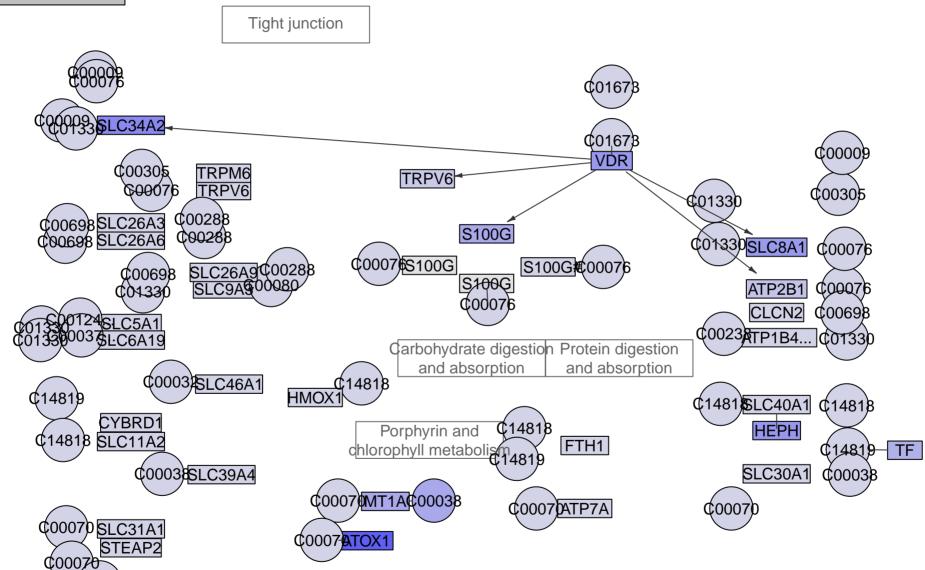


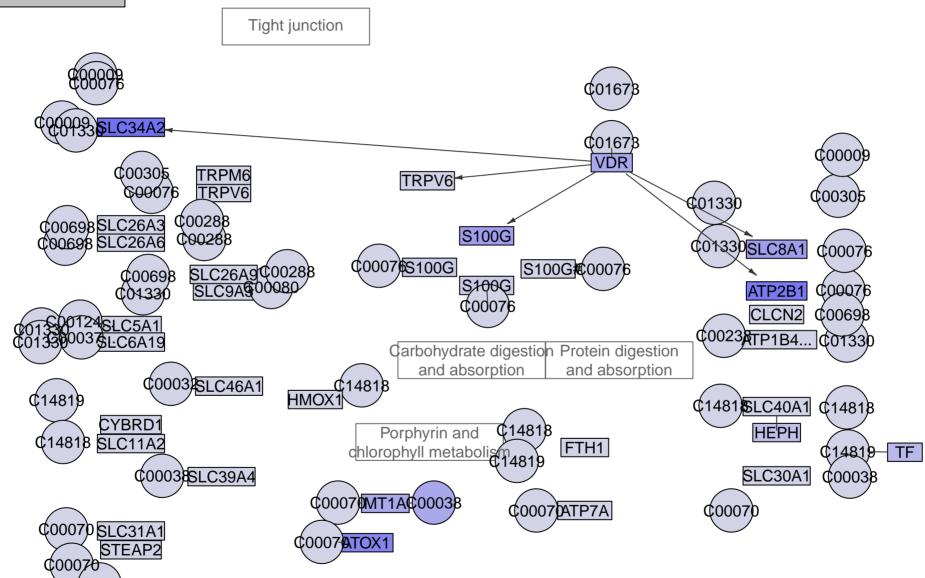
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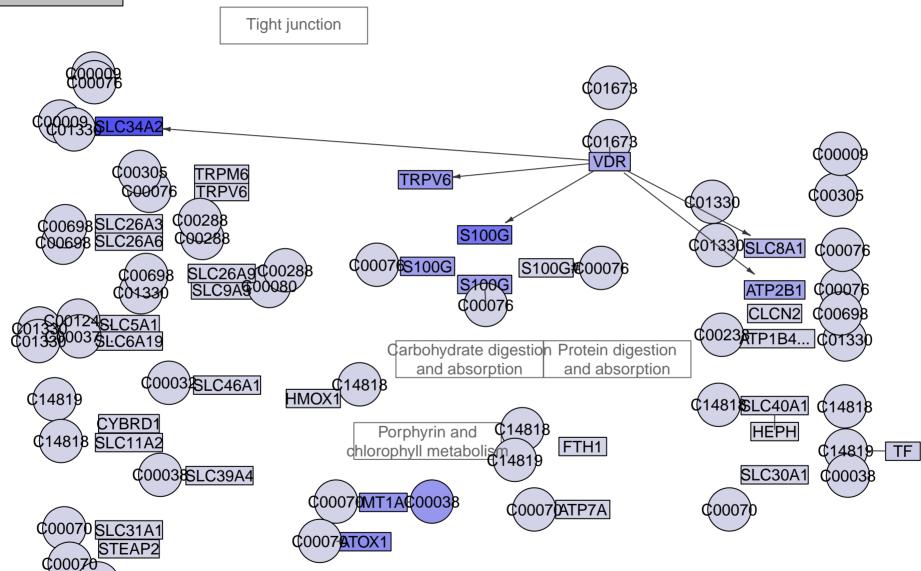


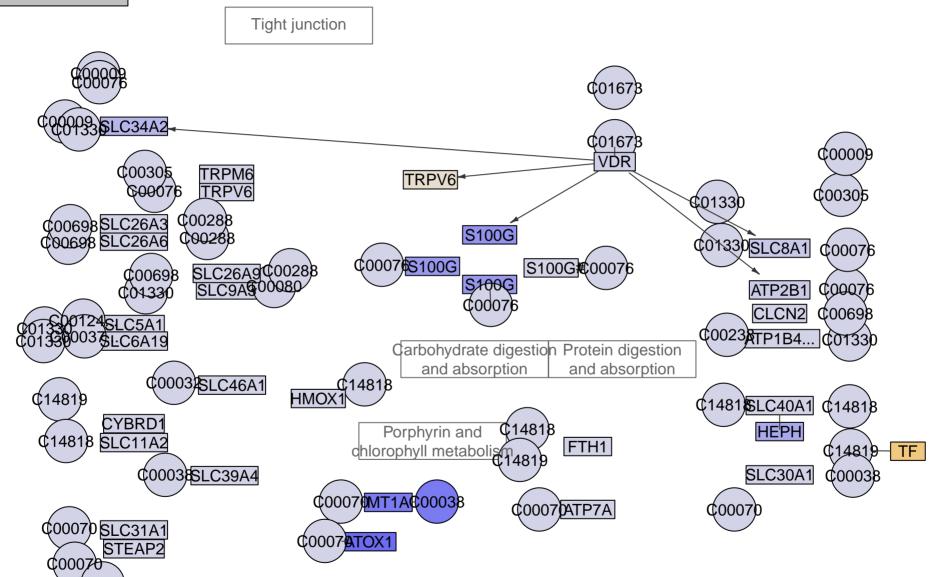


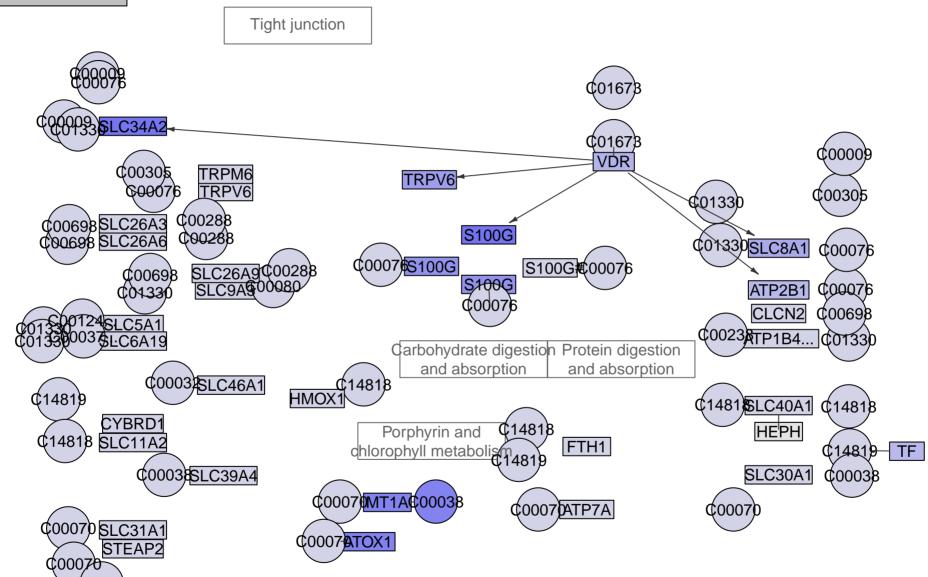


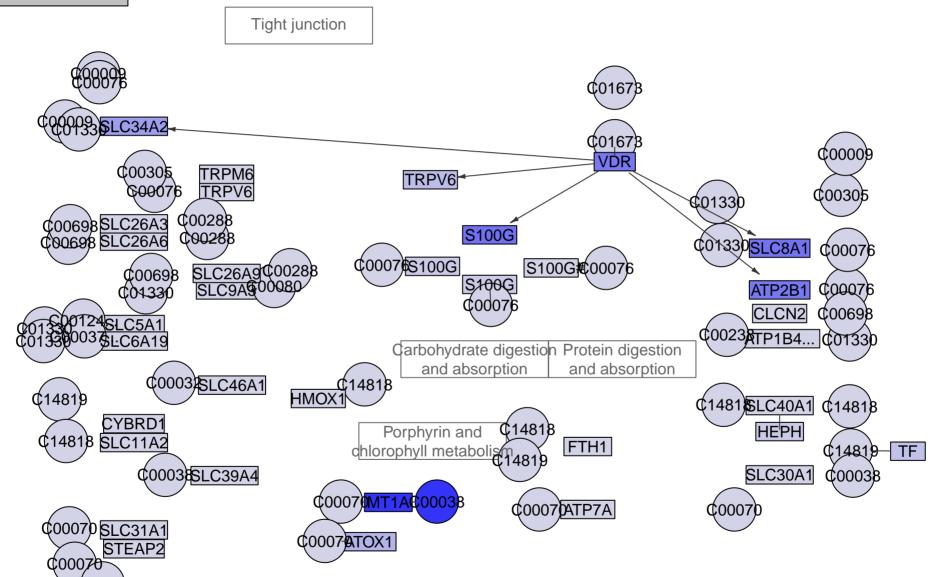


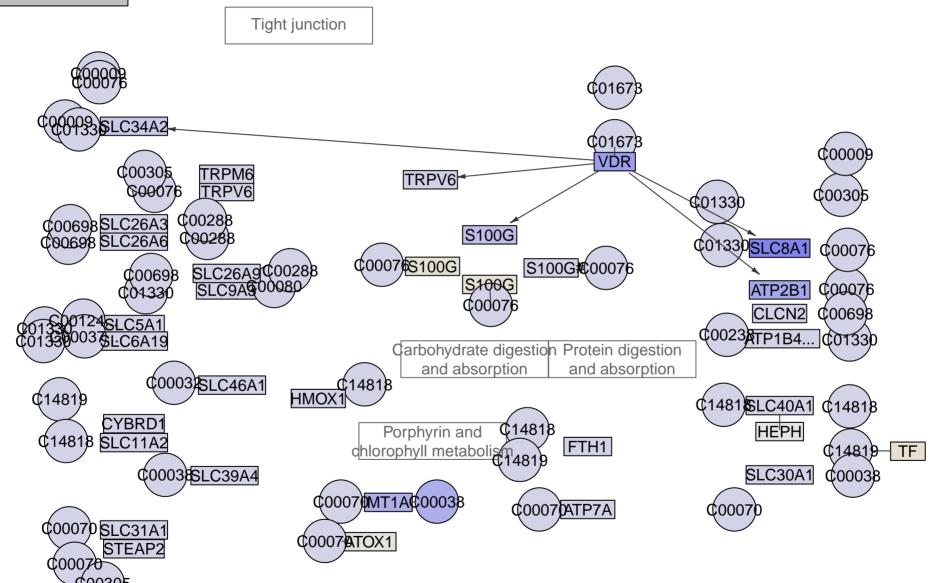


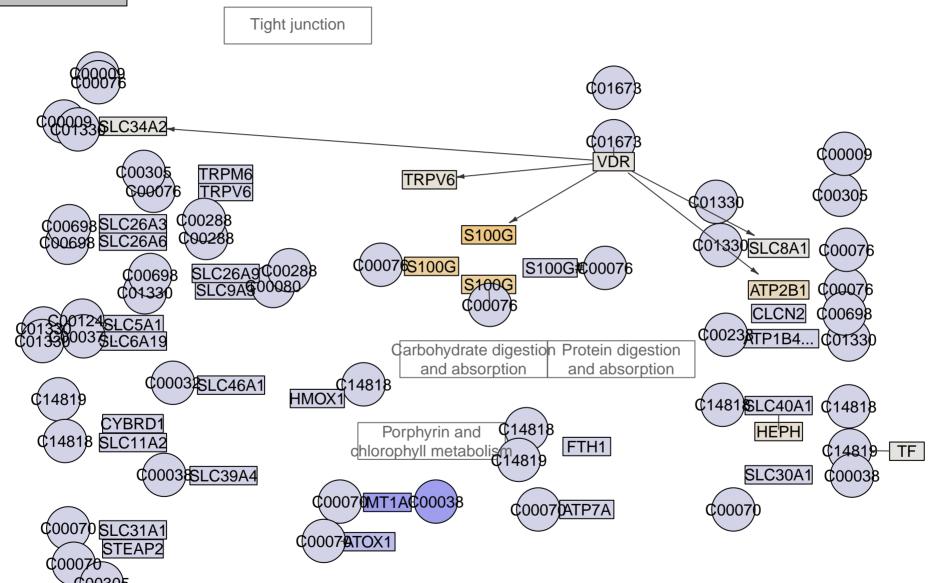


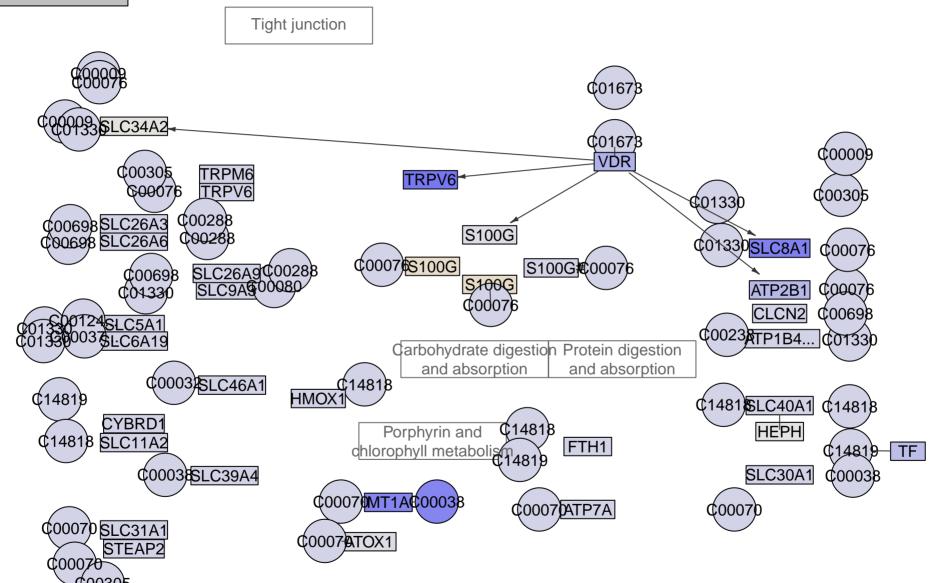


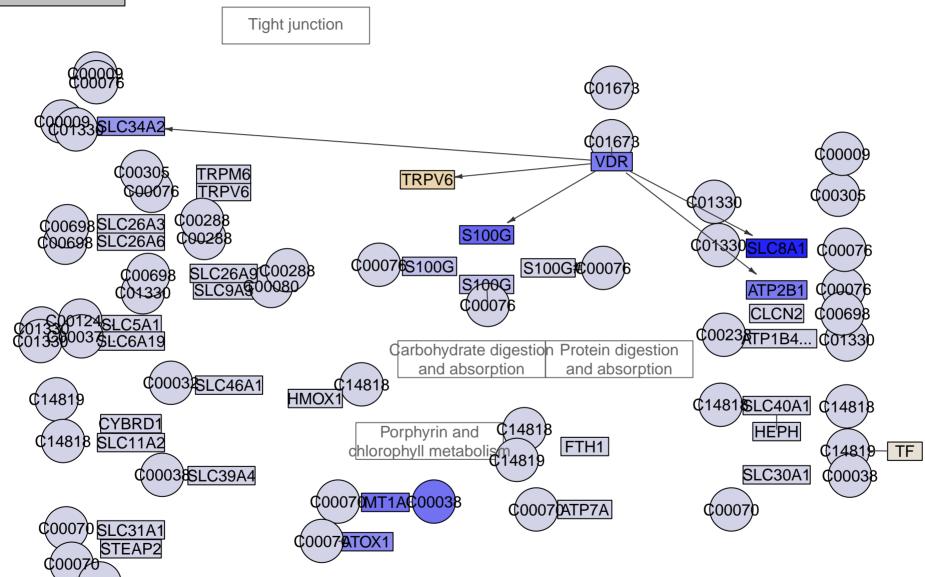


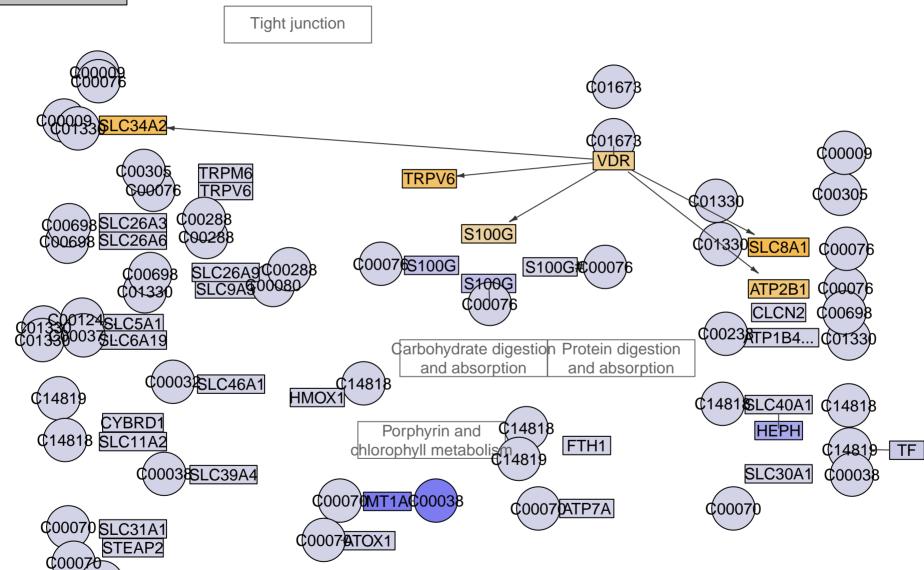


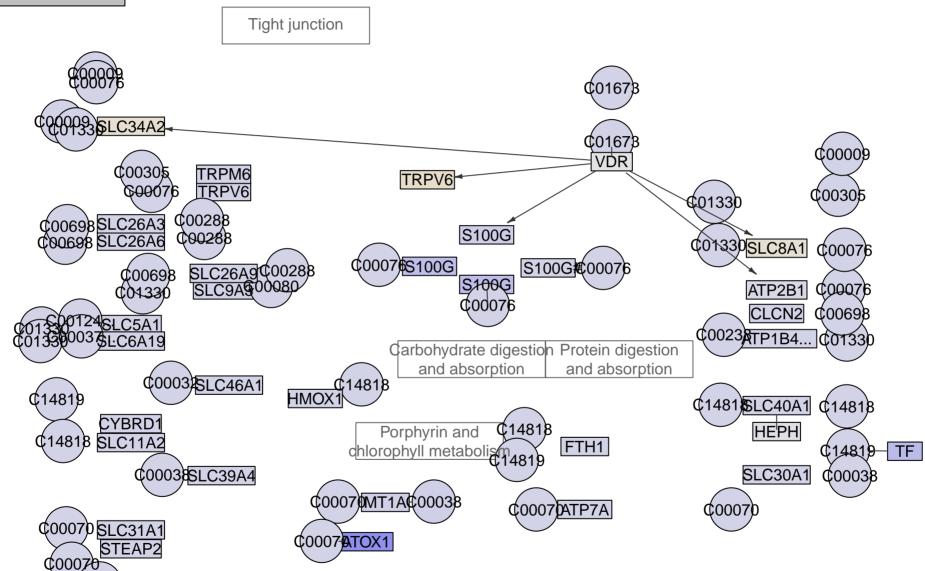


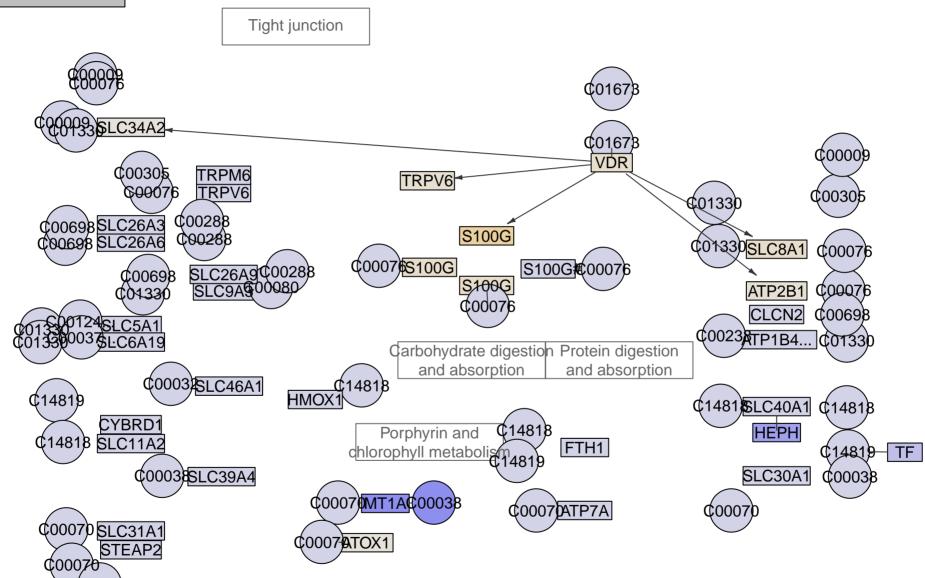


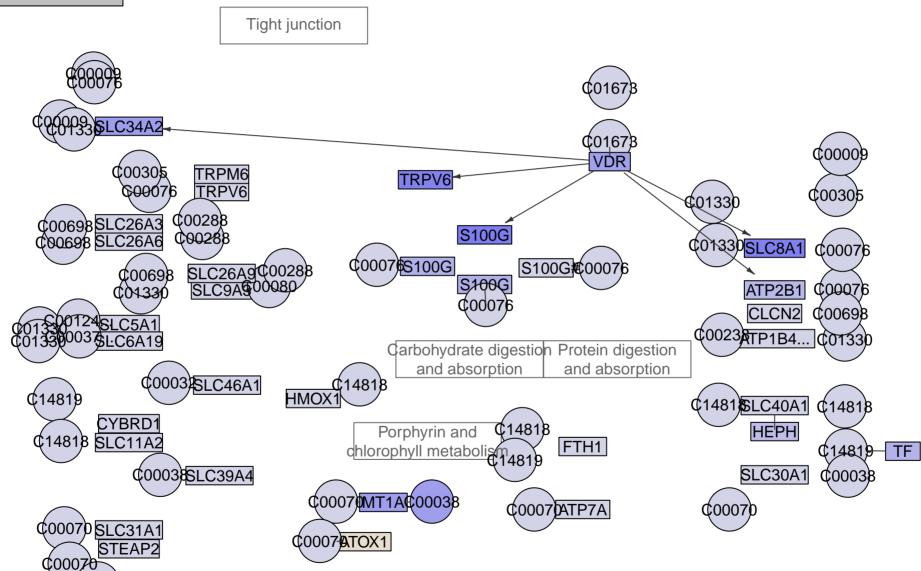


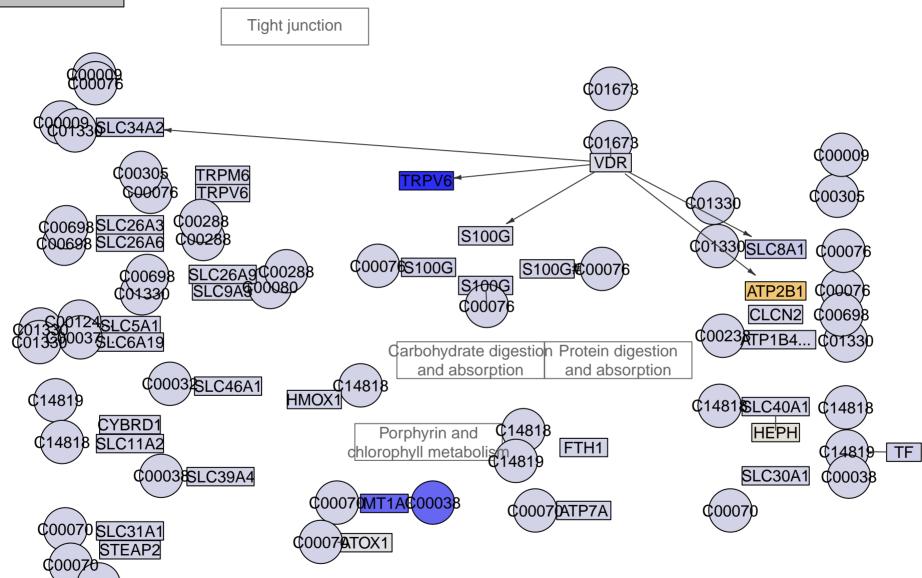


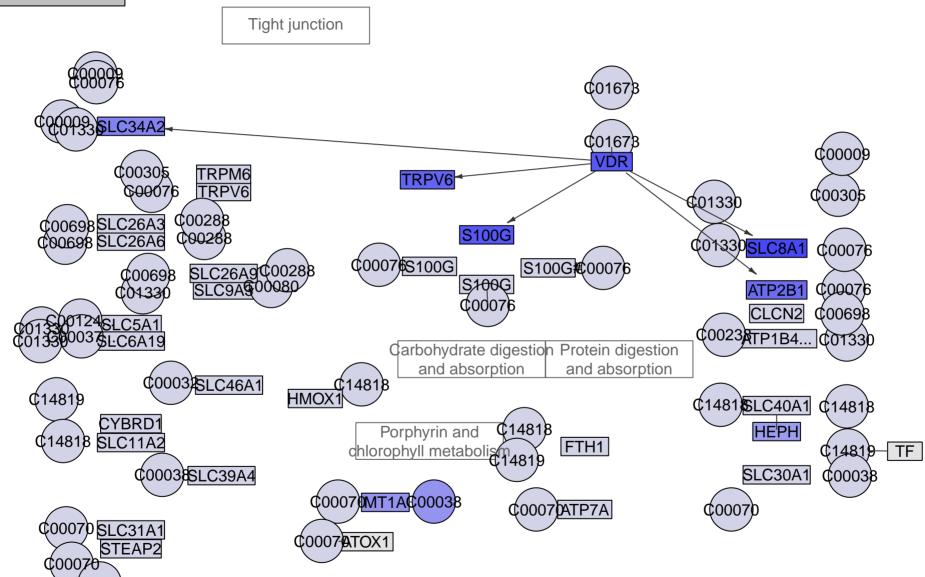


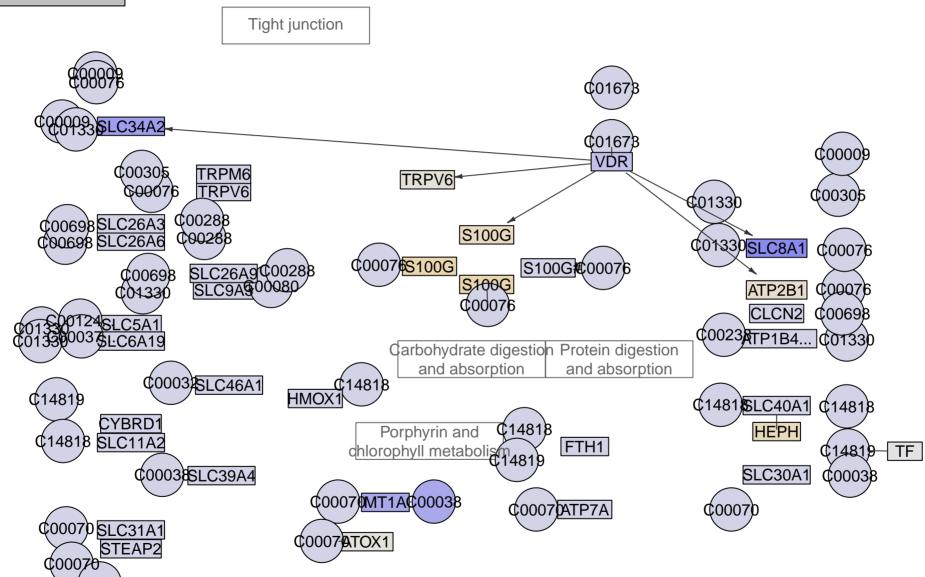


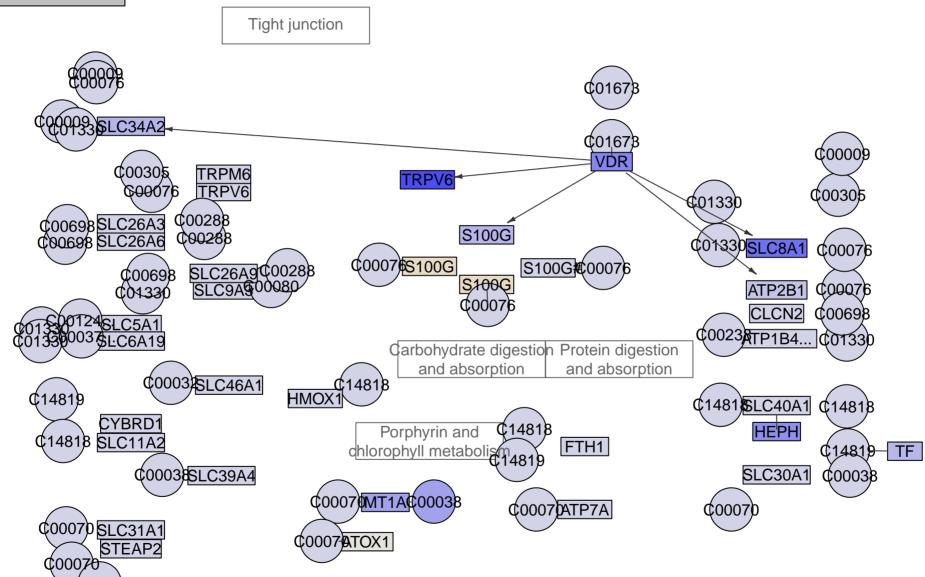


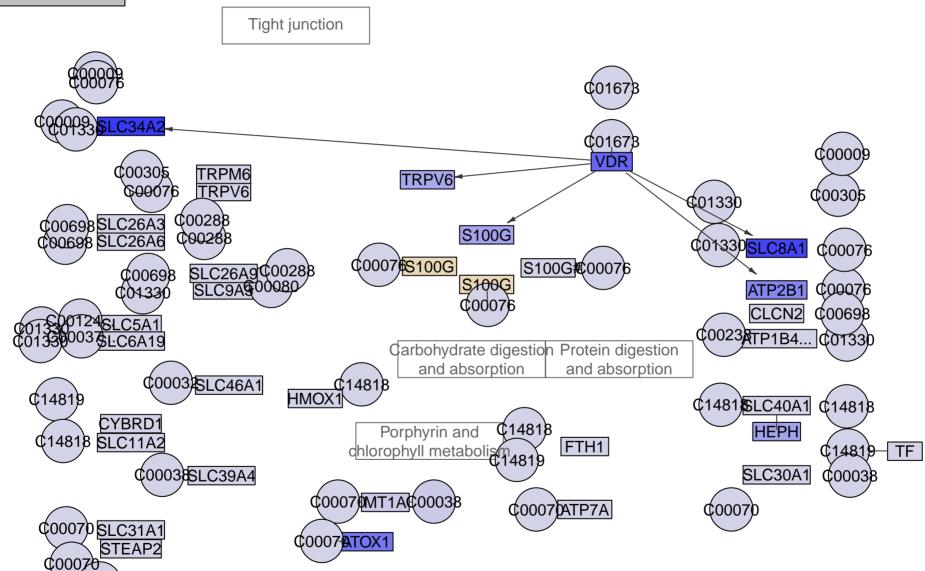


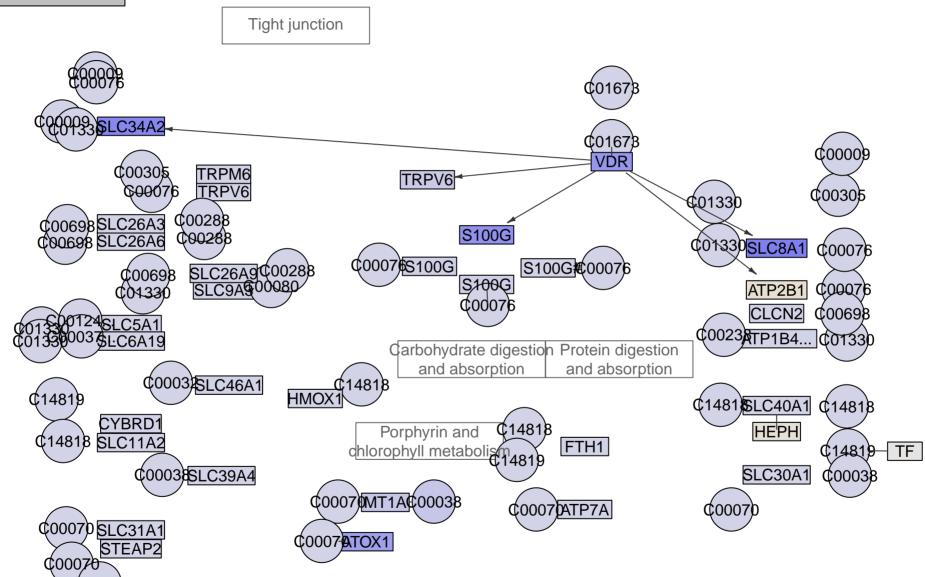


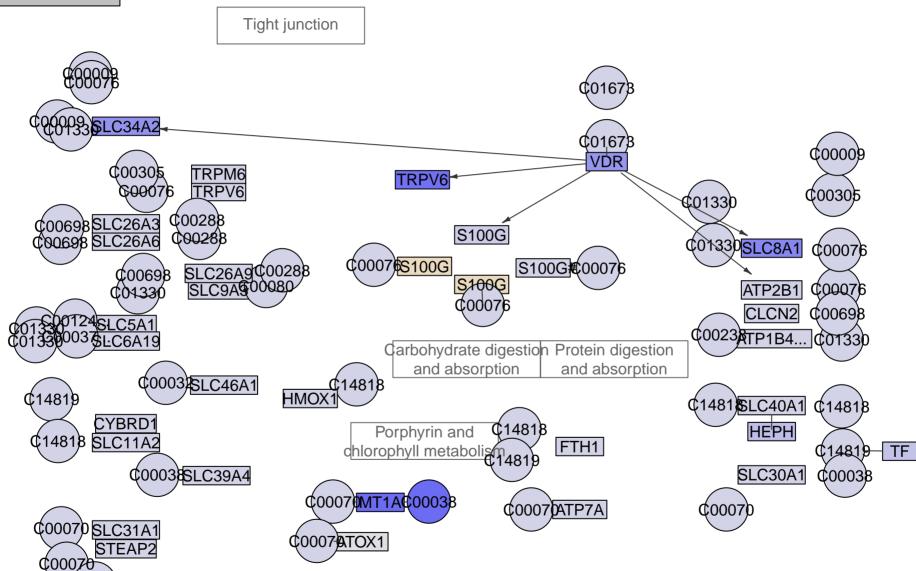


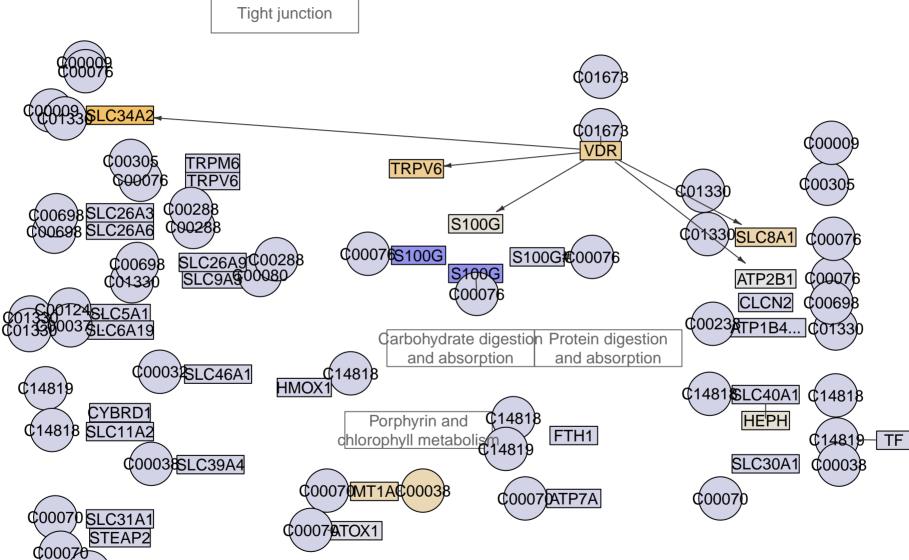


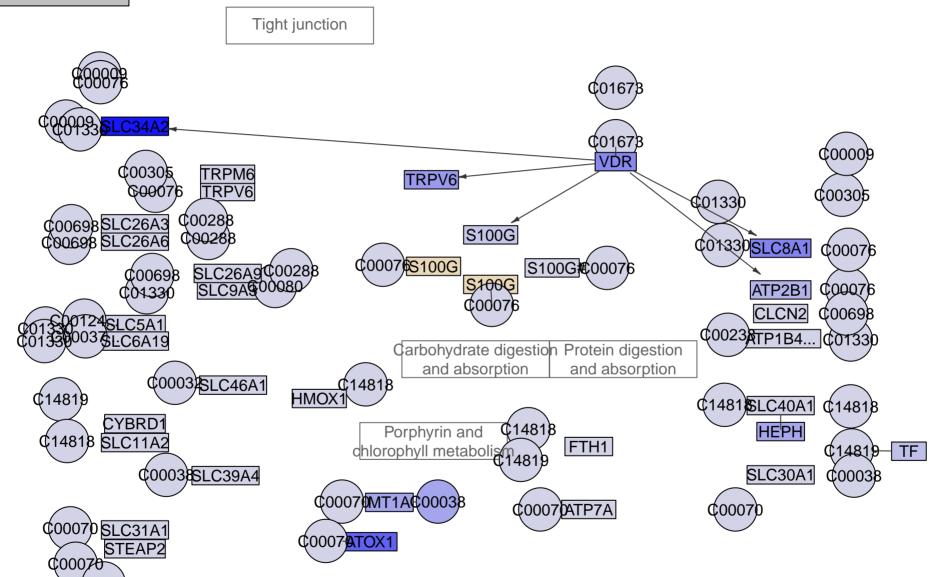






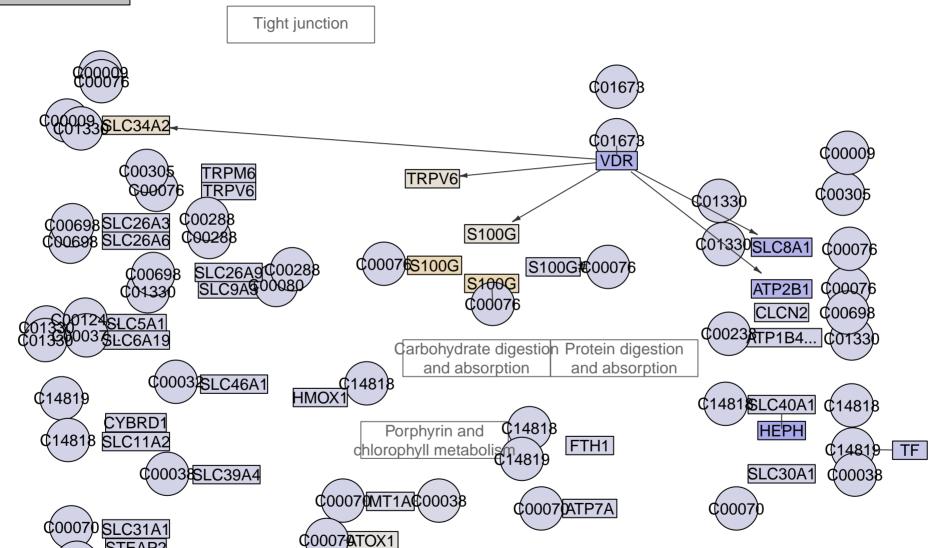




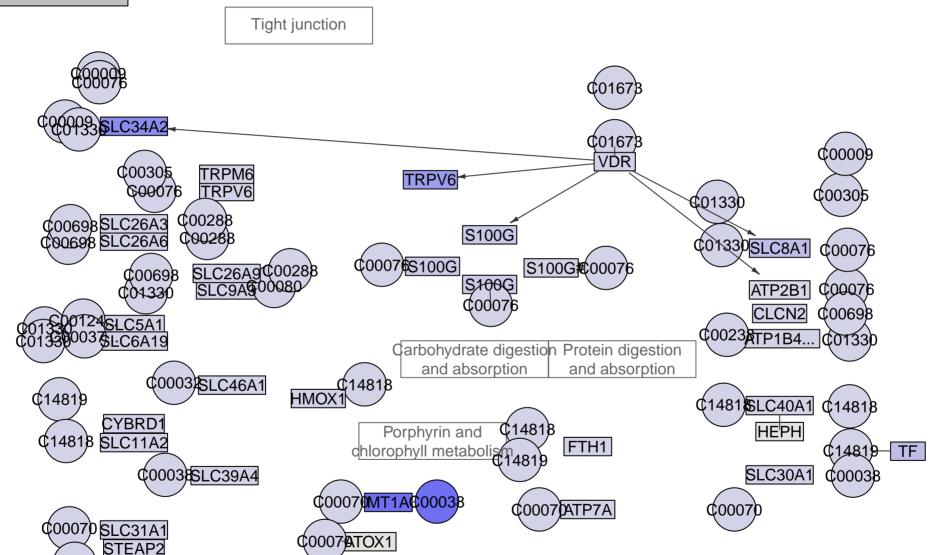


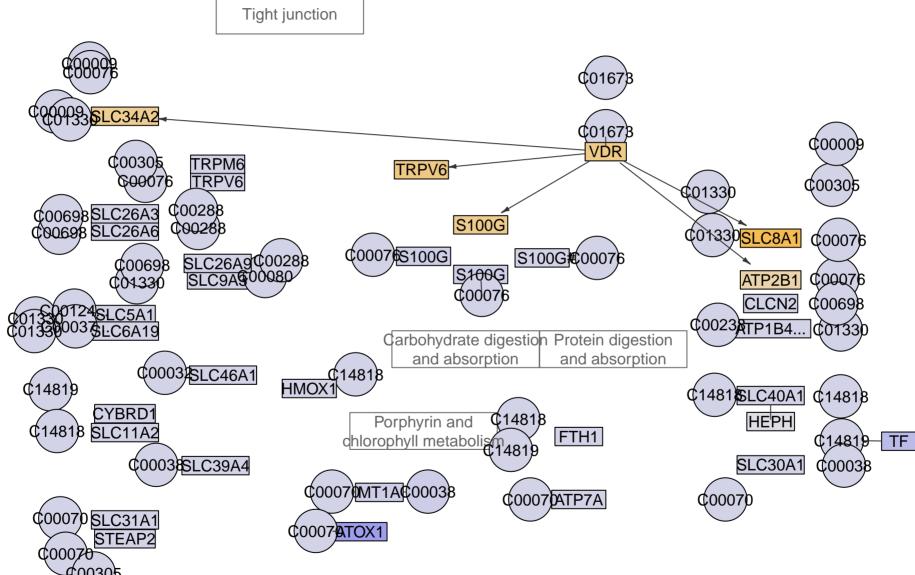
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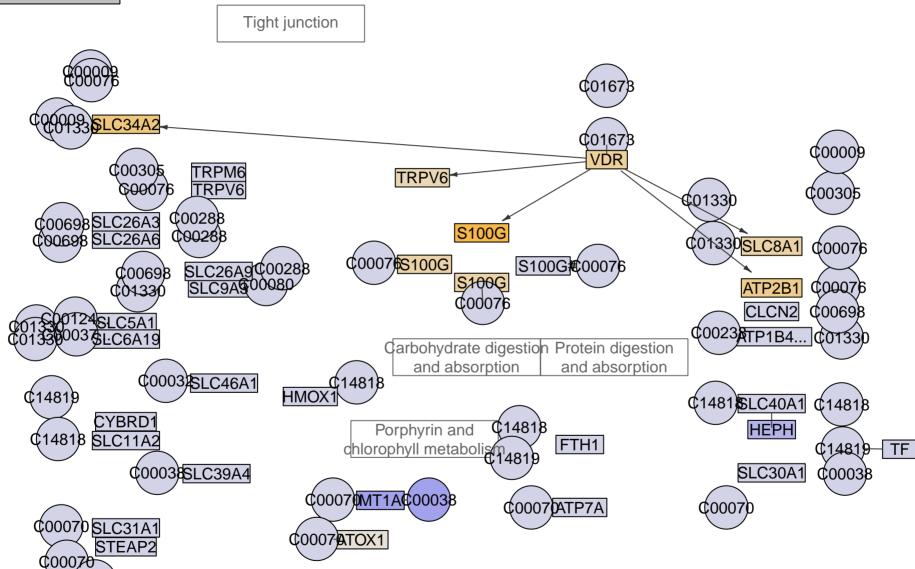
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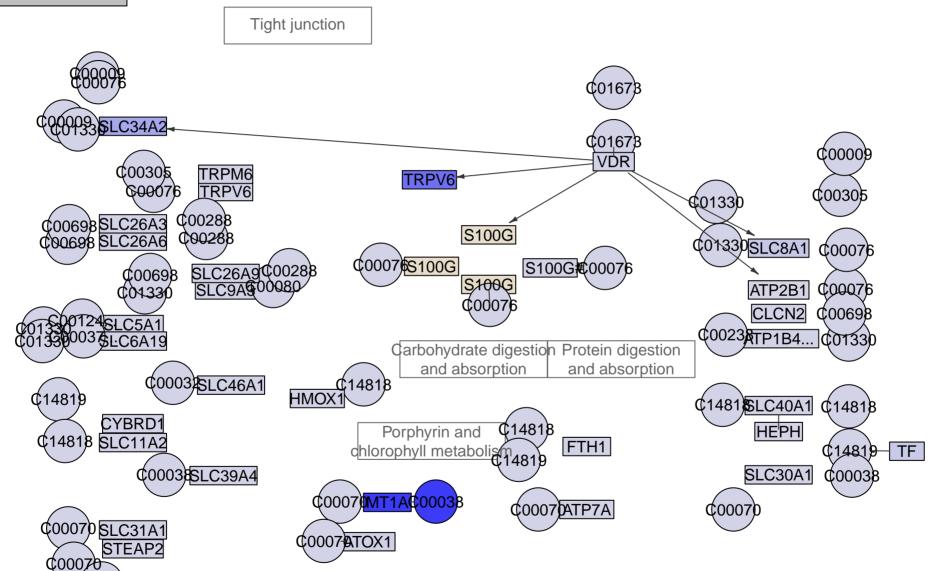


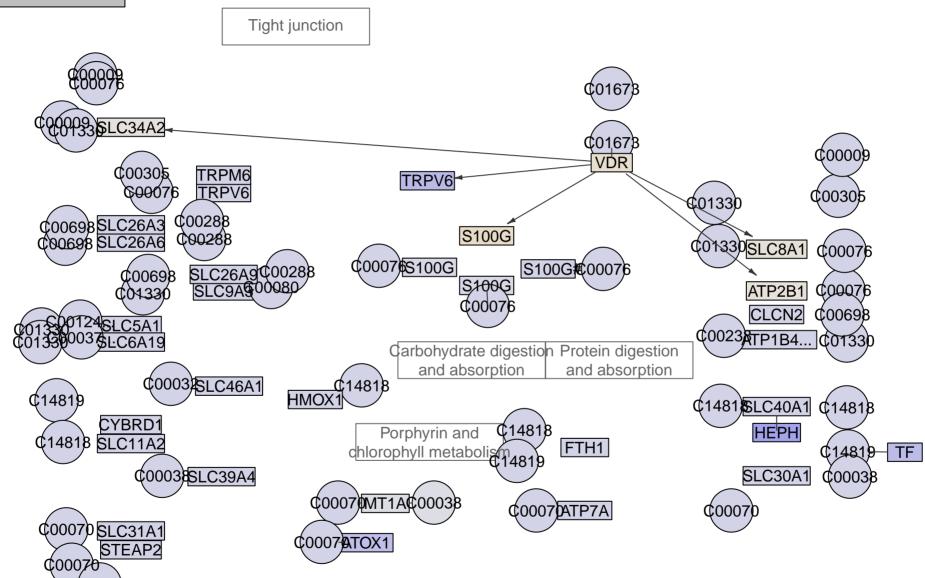
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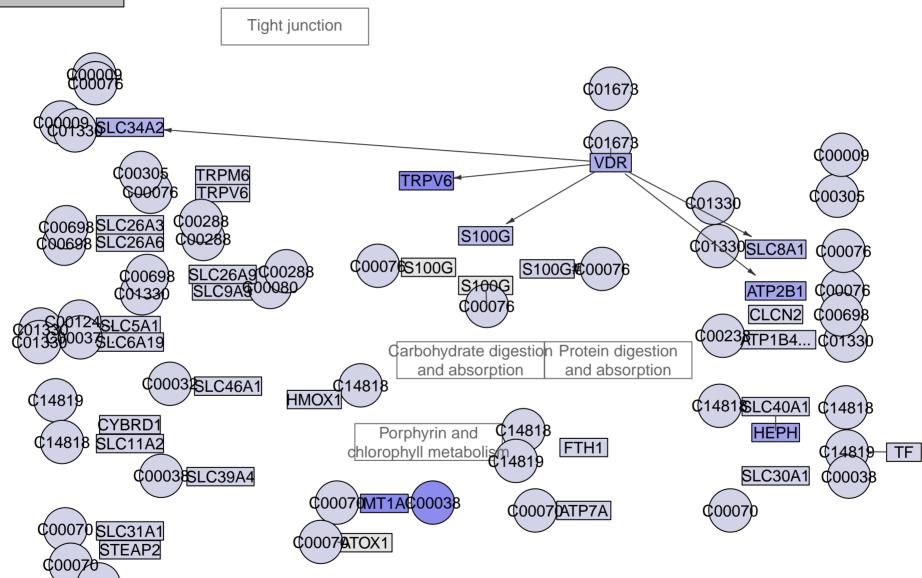




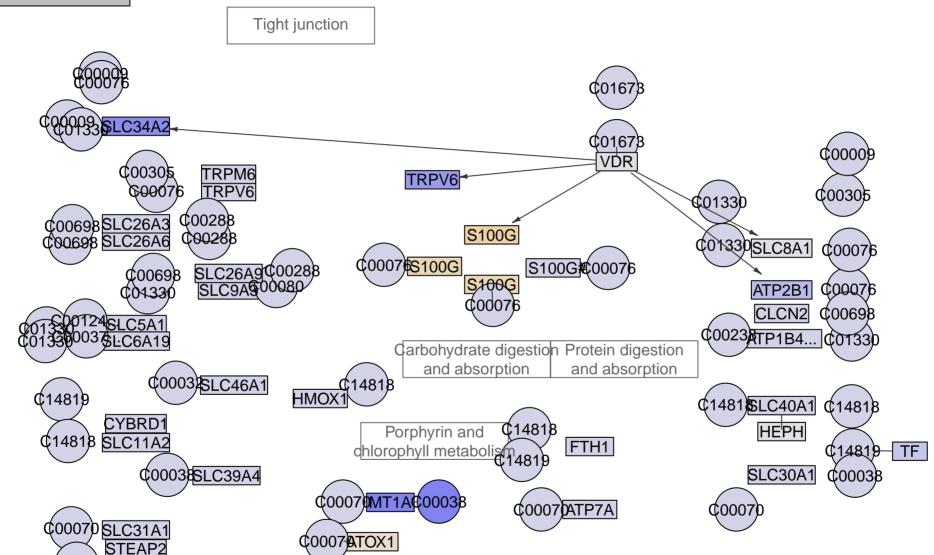


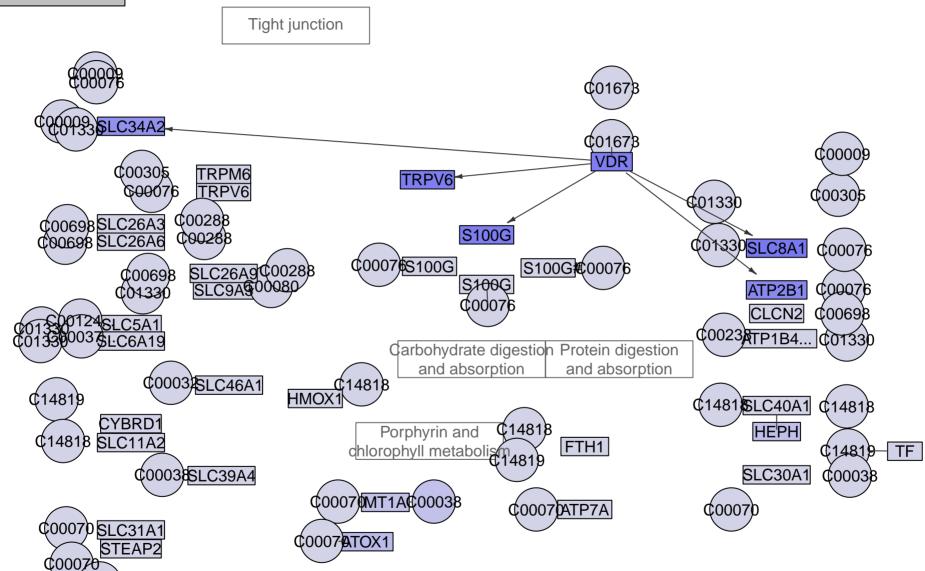


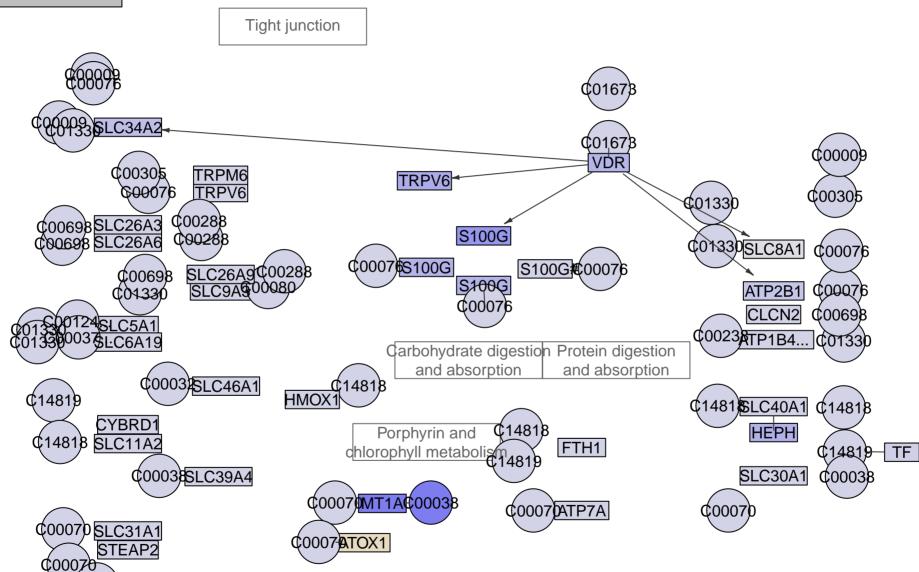


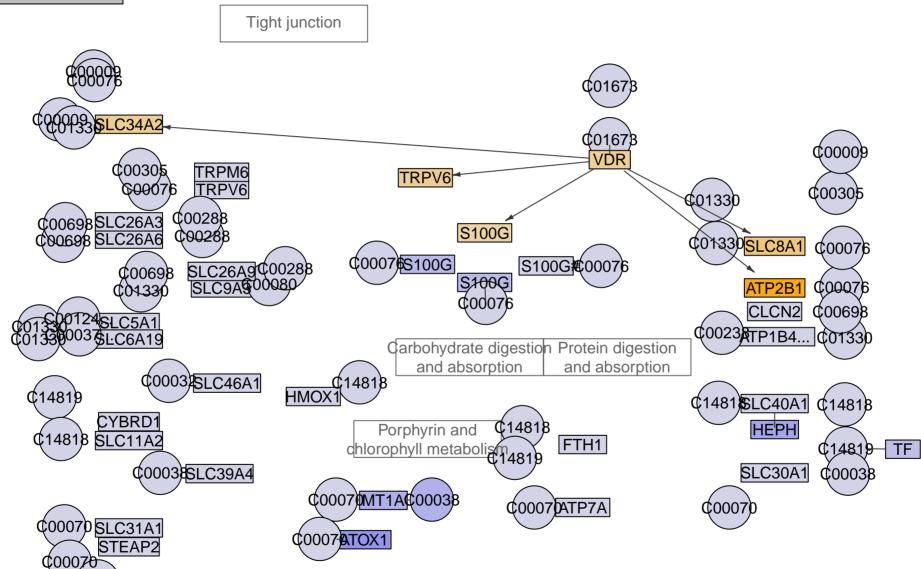


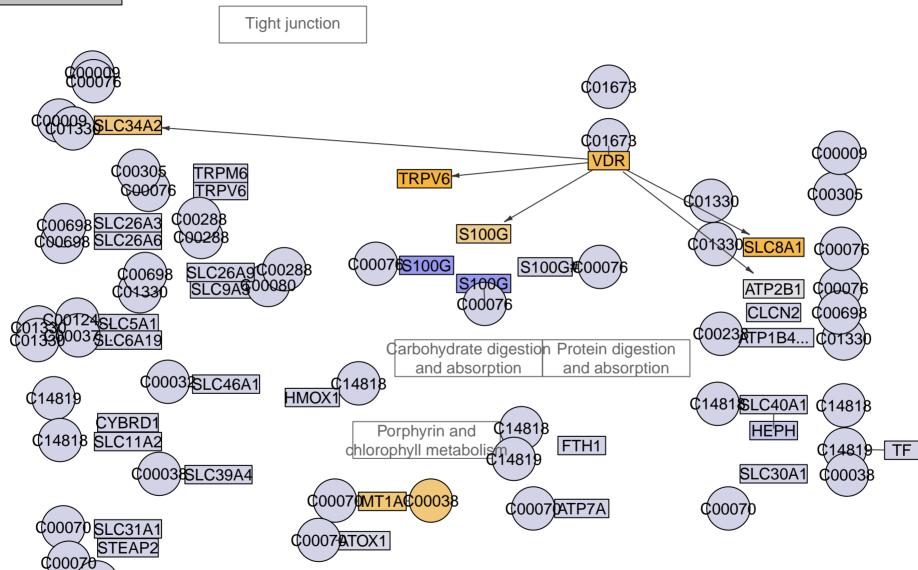
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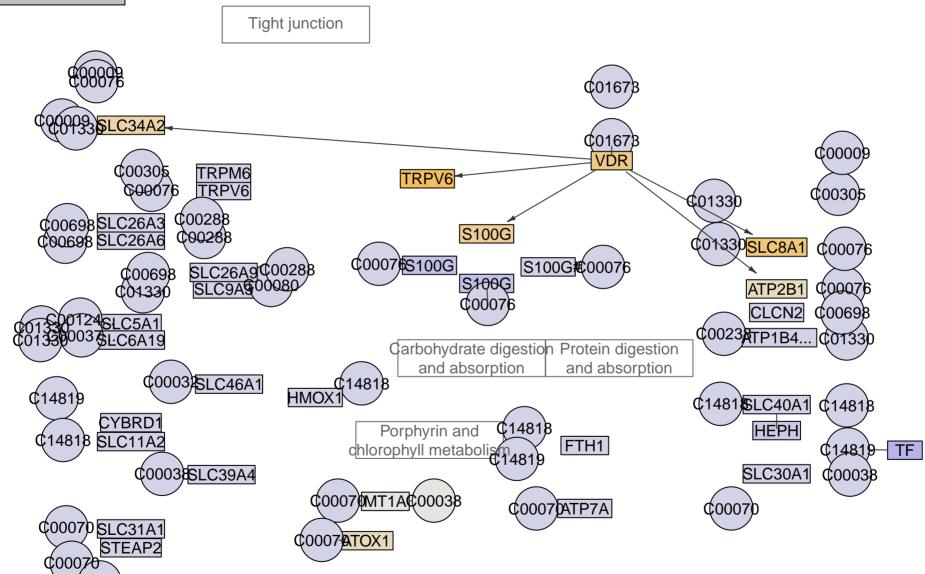


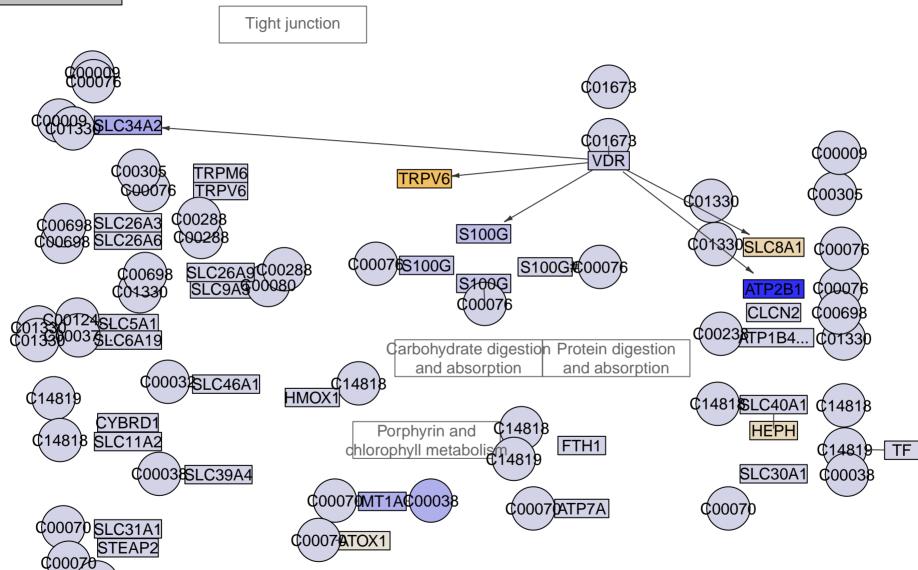






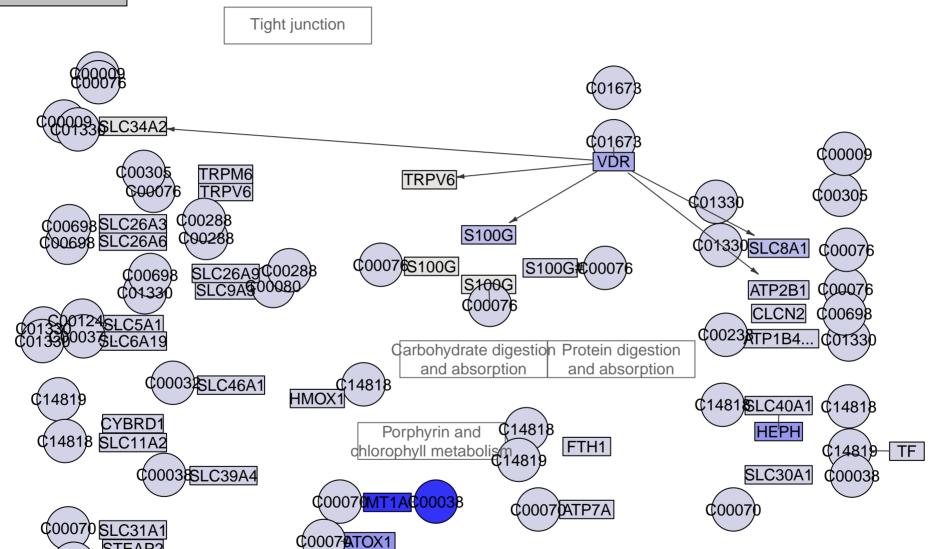


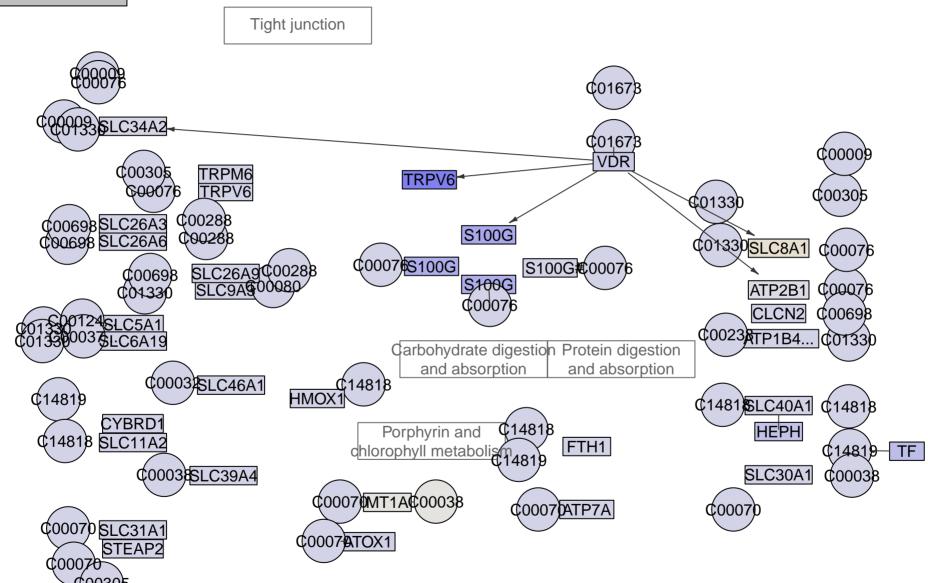


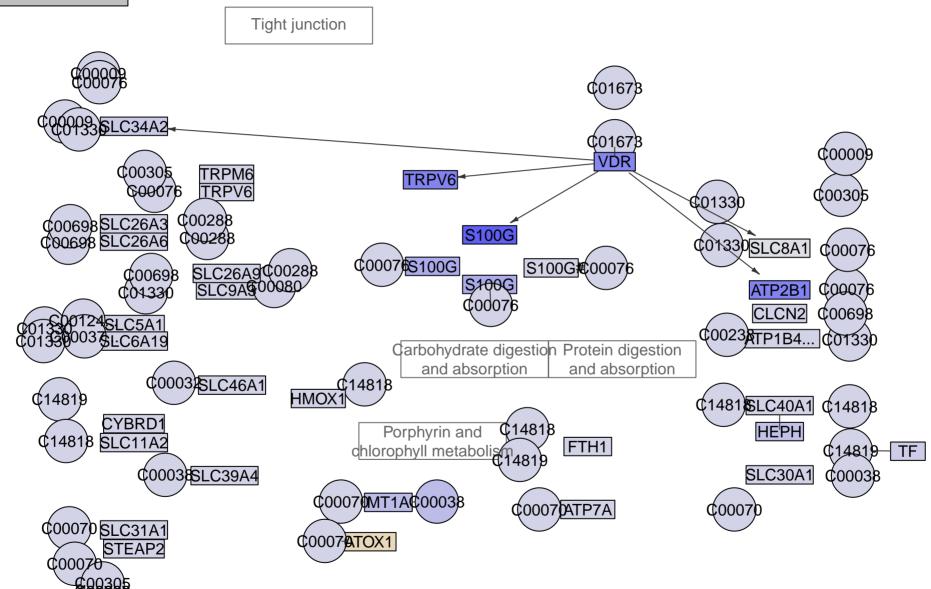


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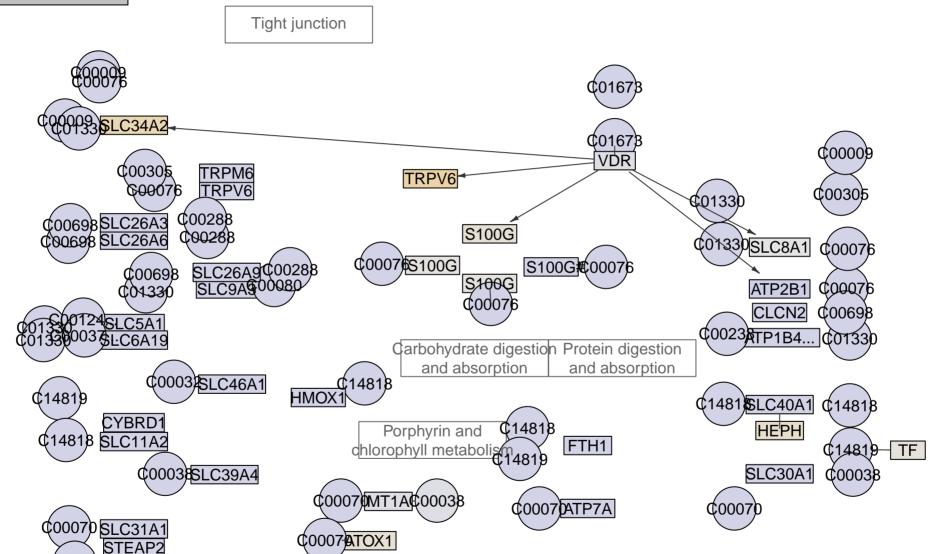
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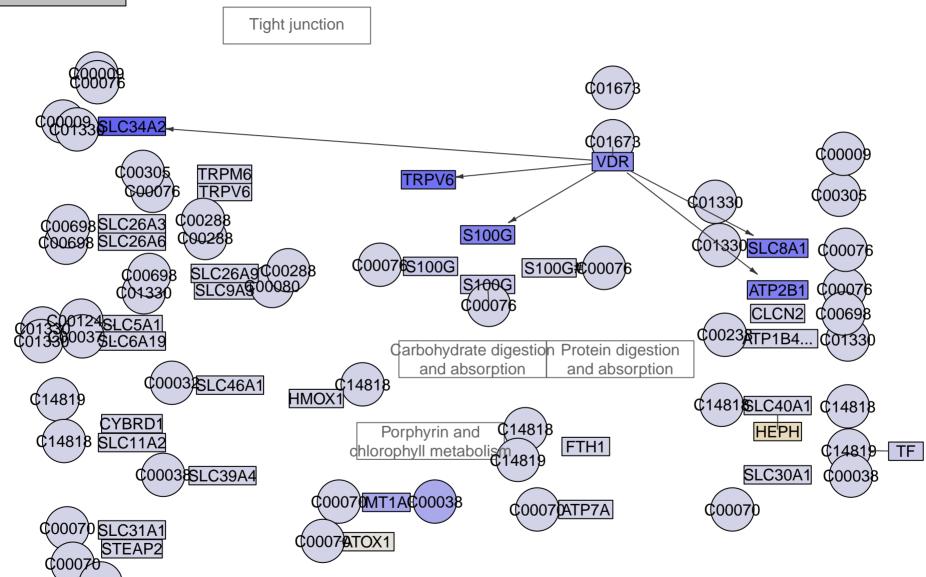


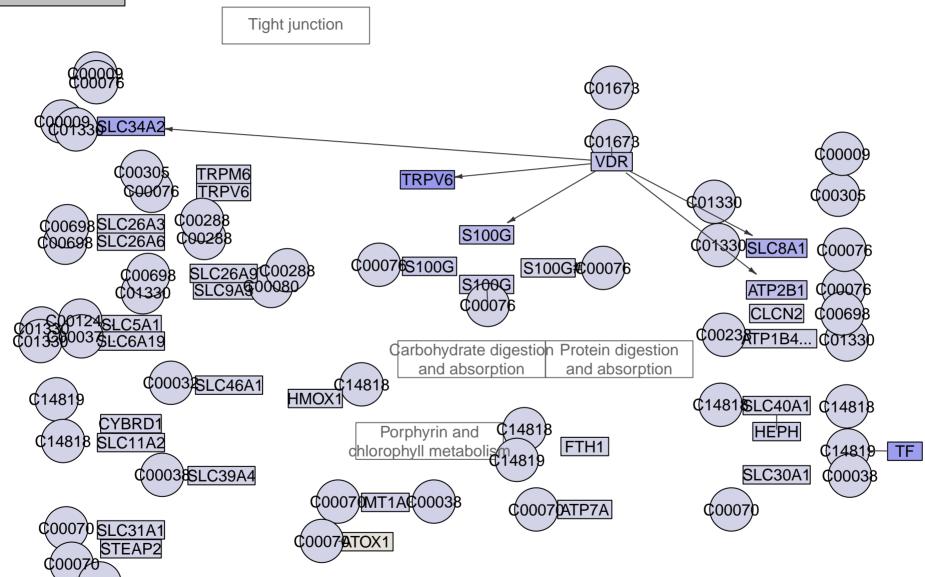


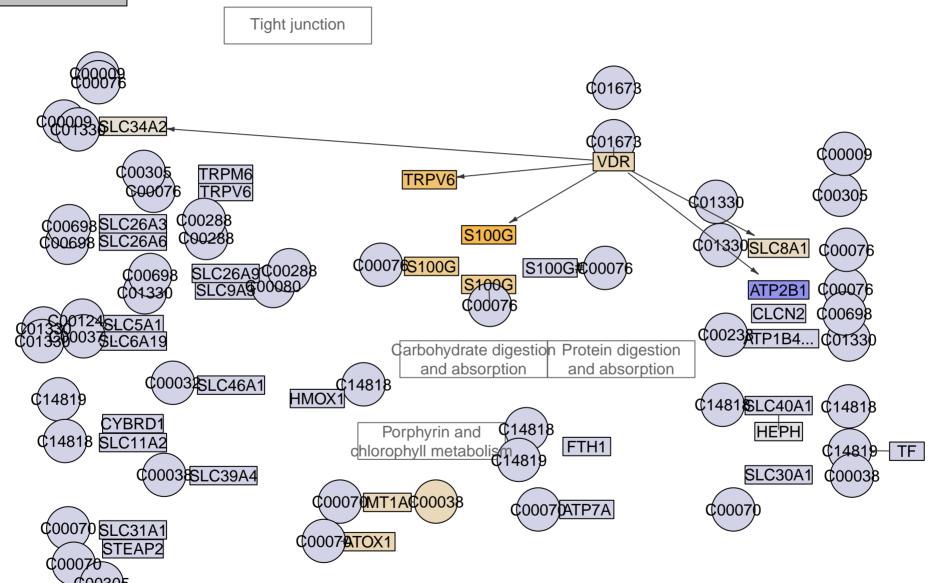


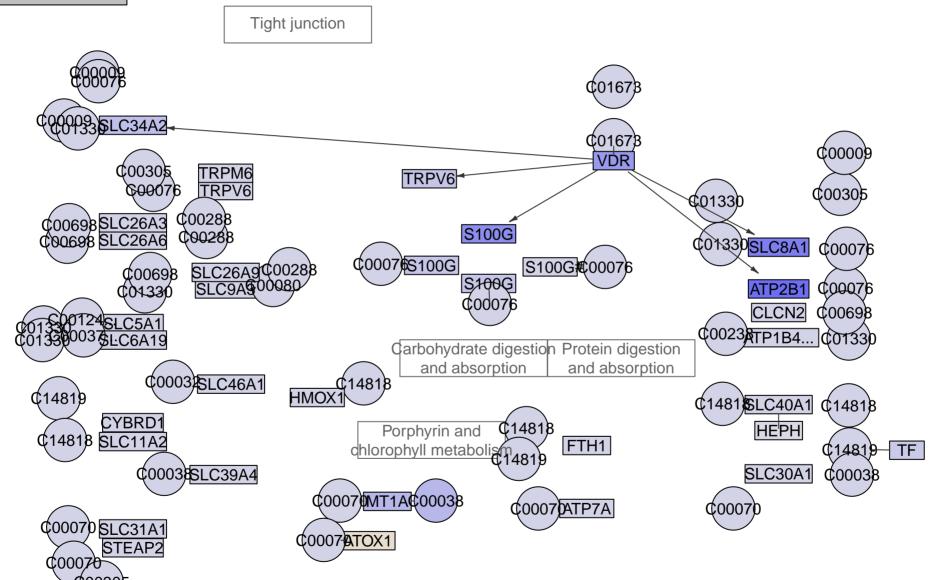
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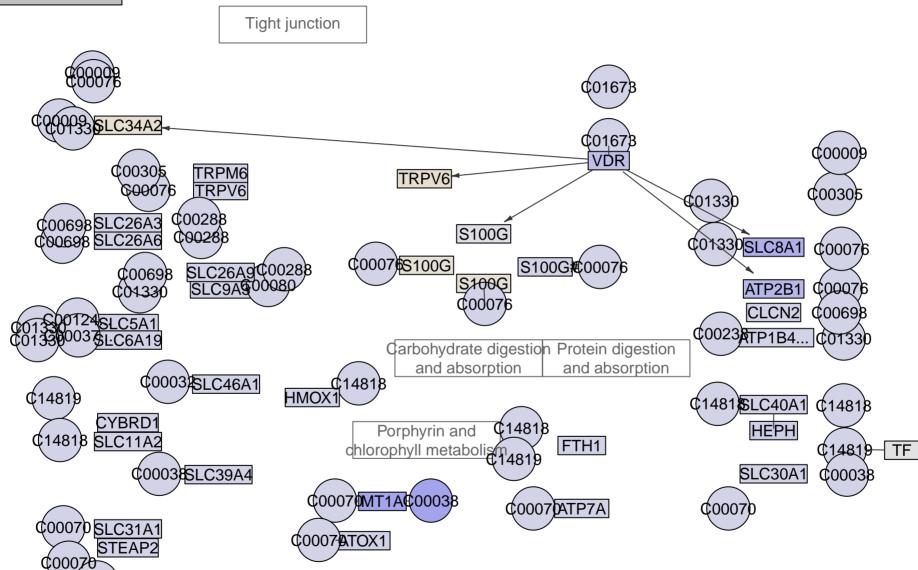




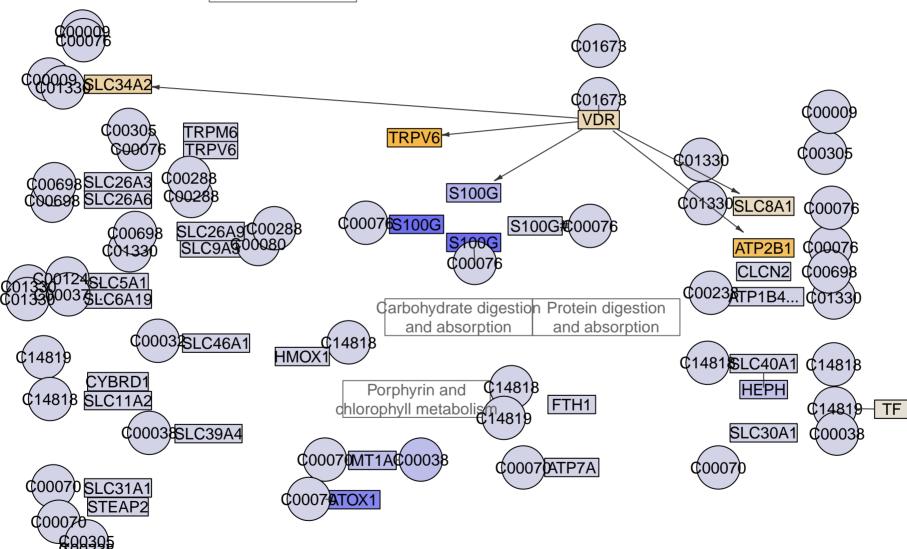


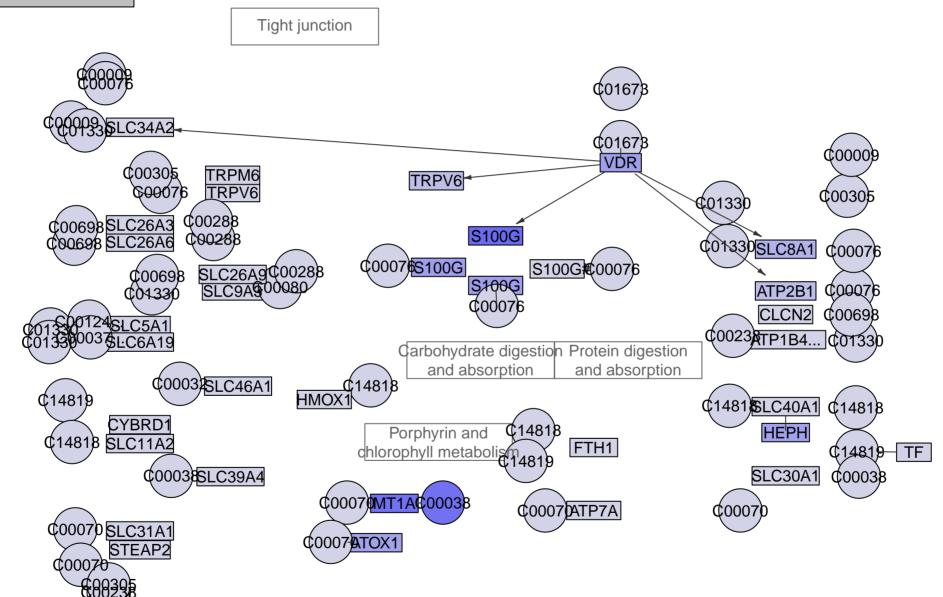


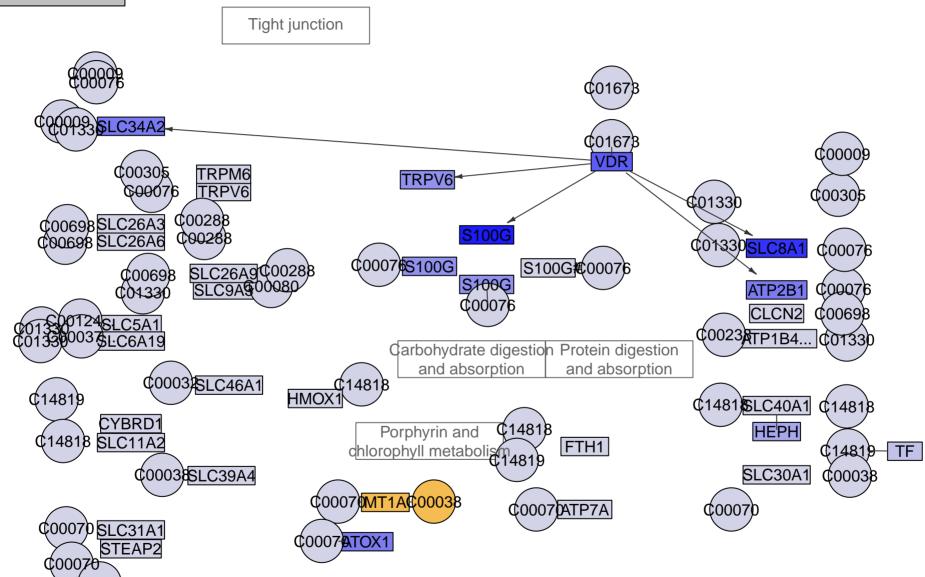


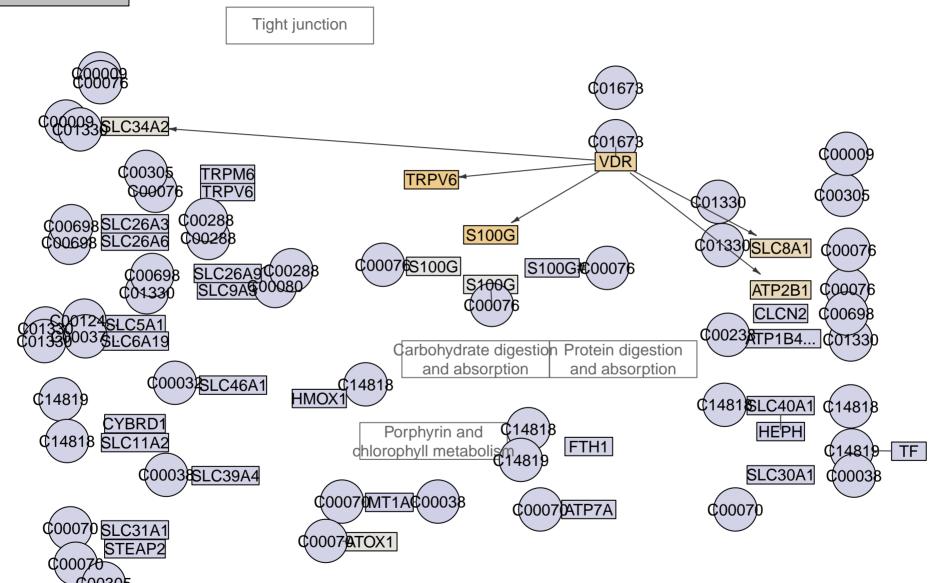


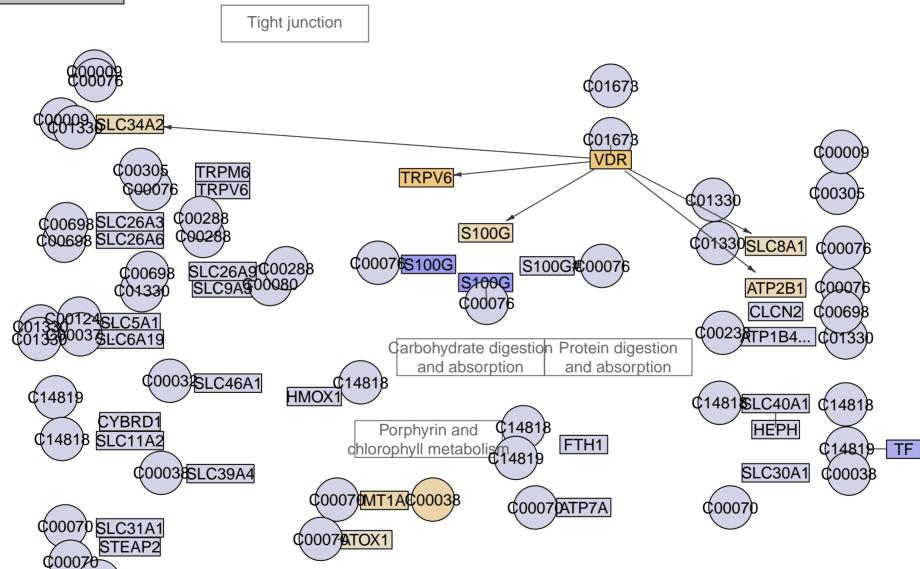
Tight junction



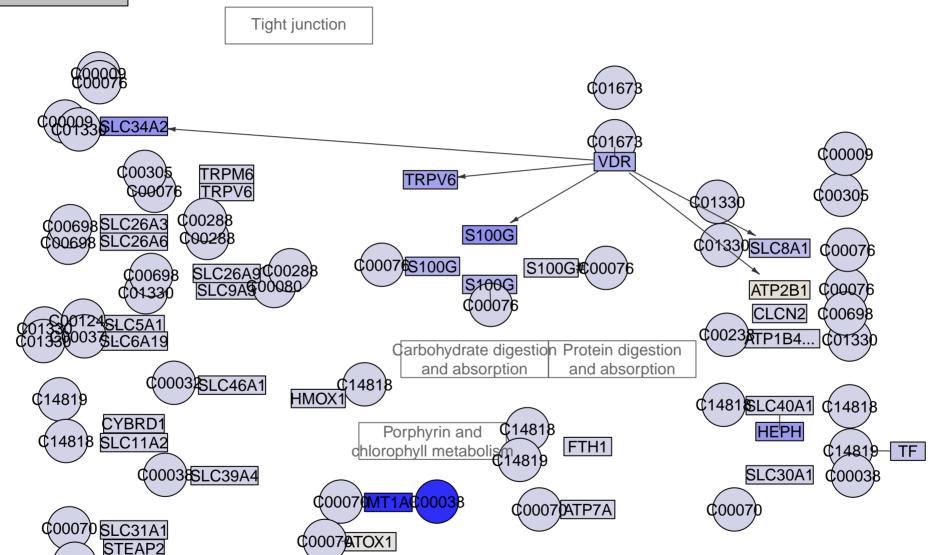




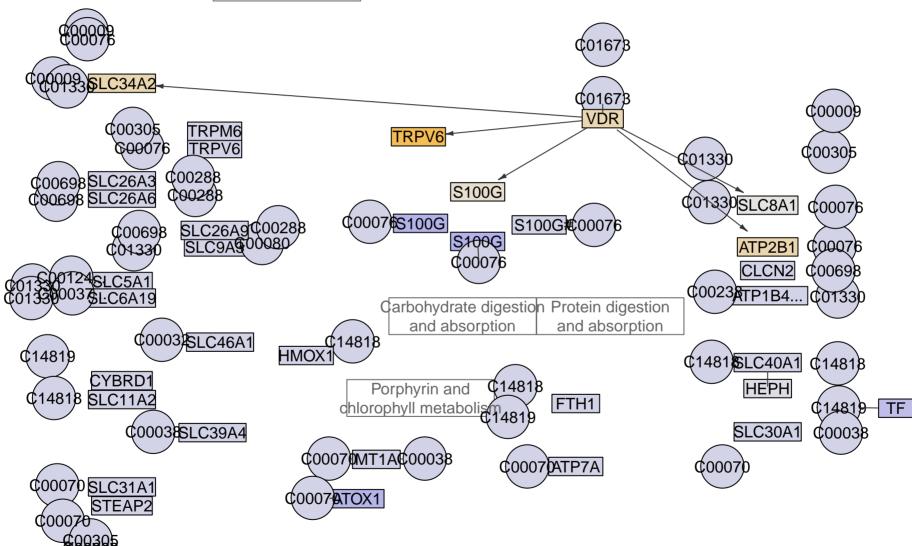


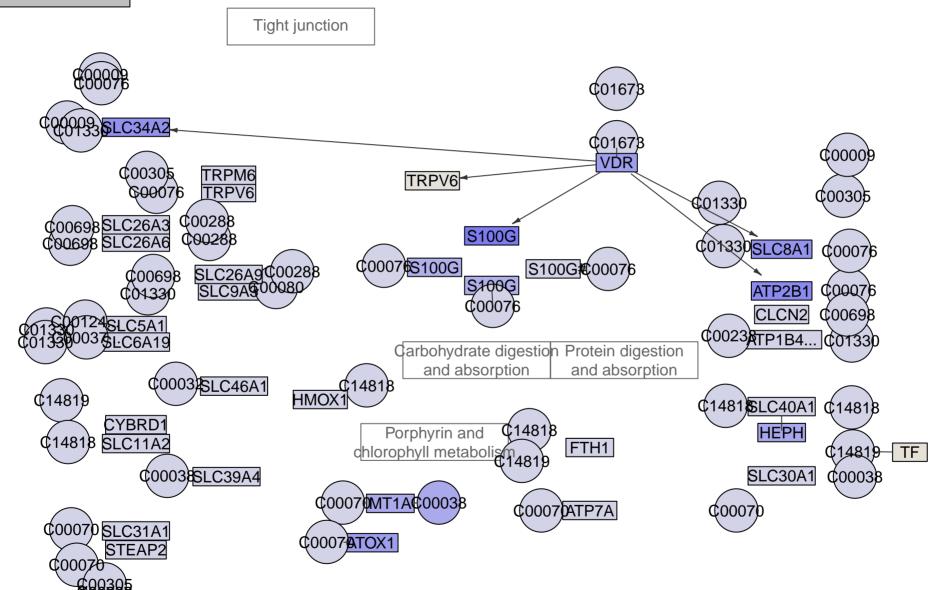


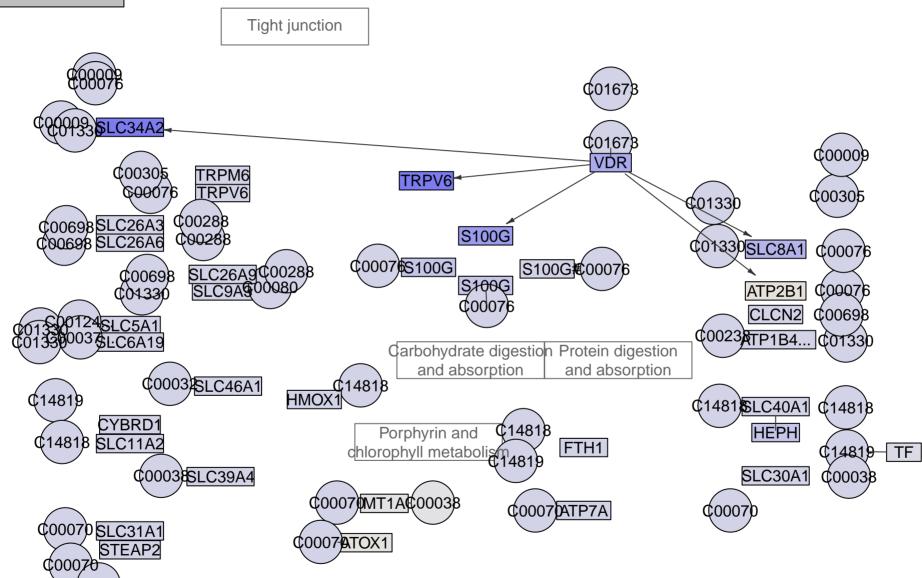
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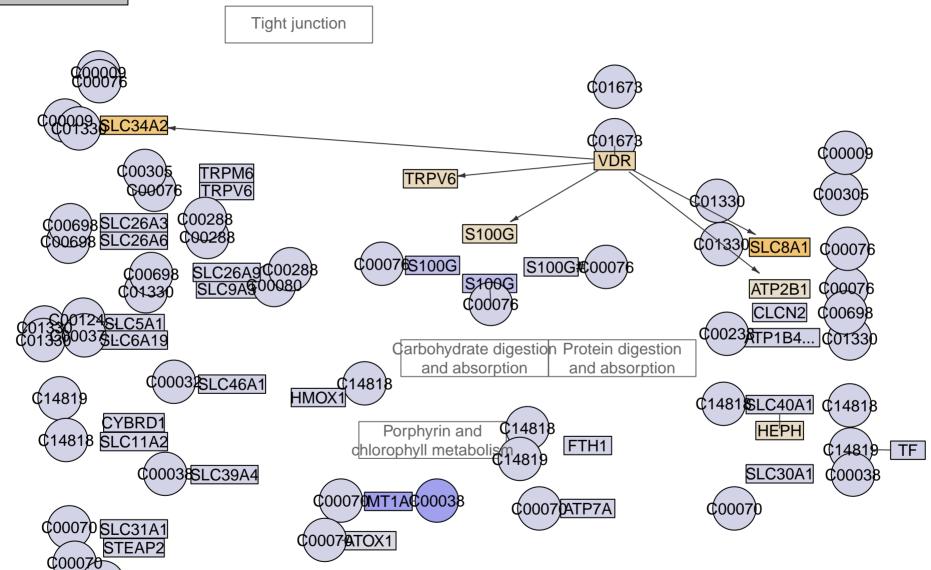


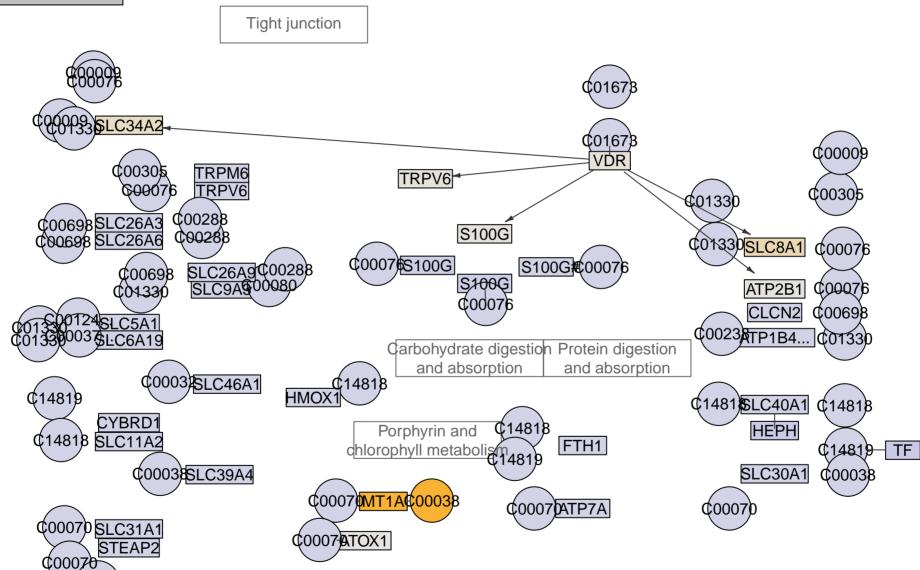
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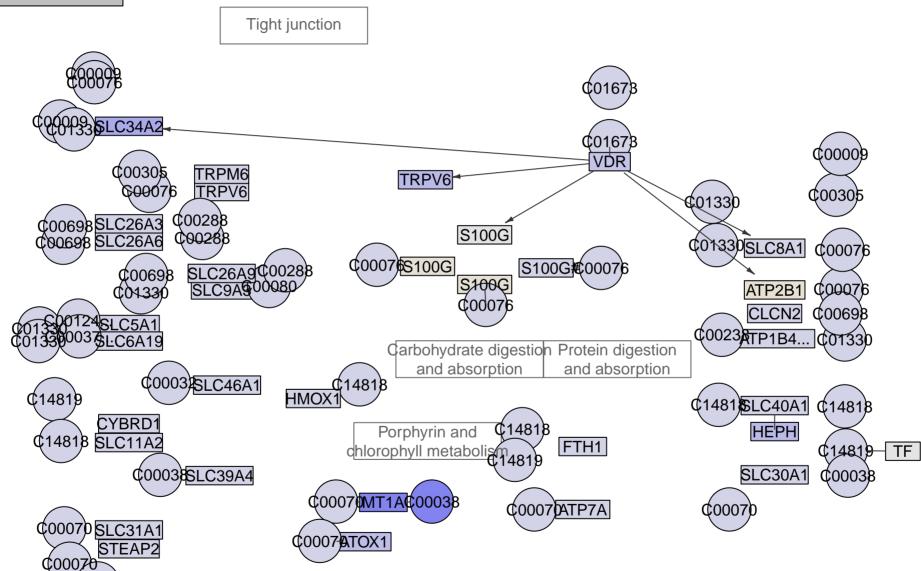


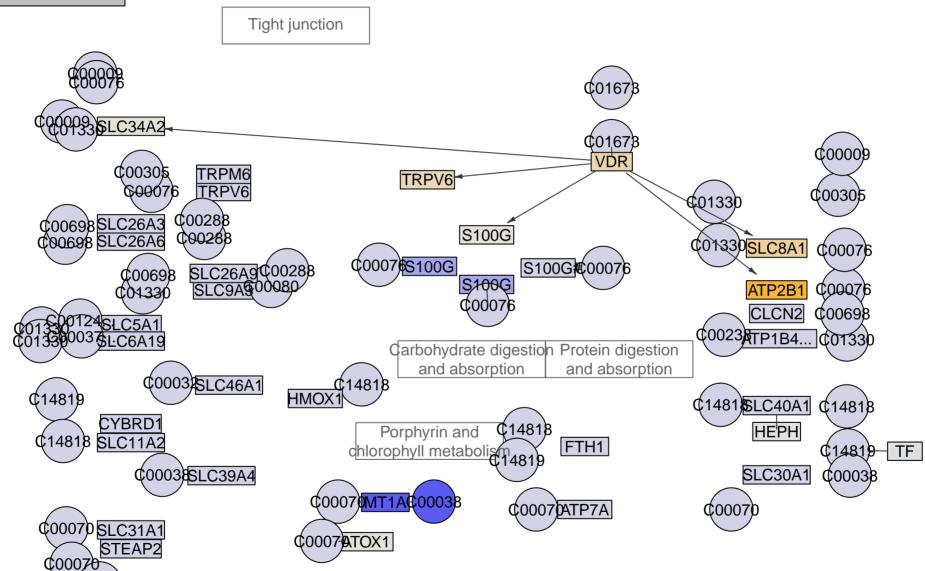


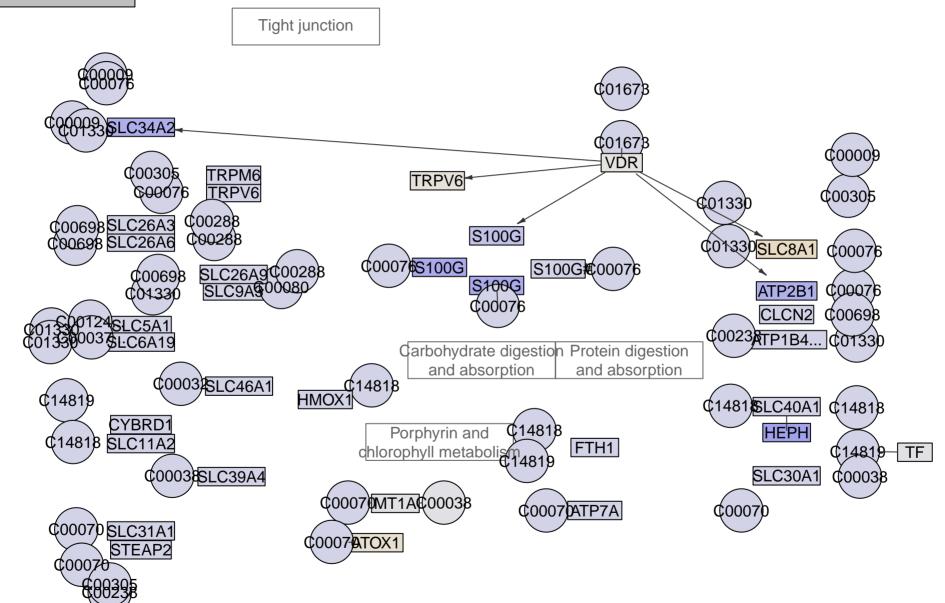






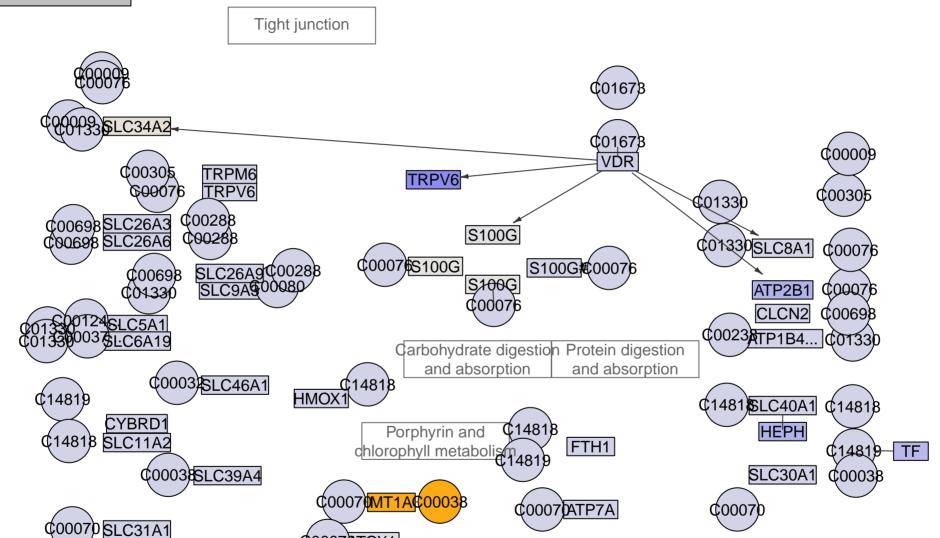




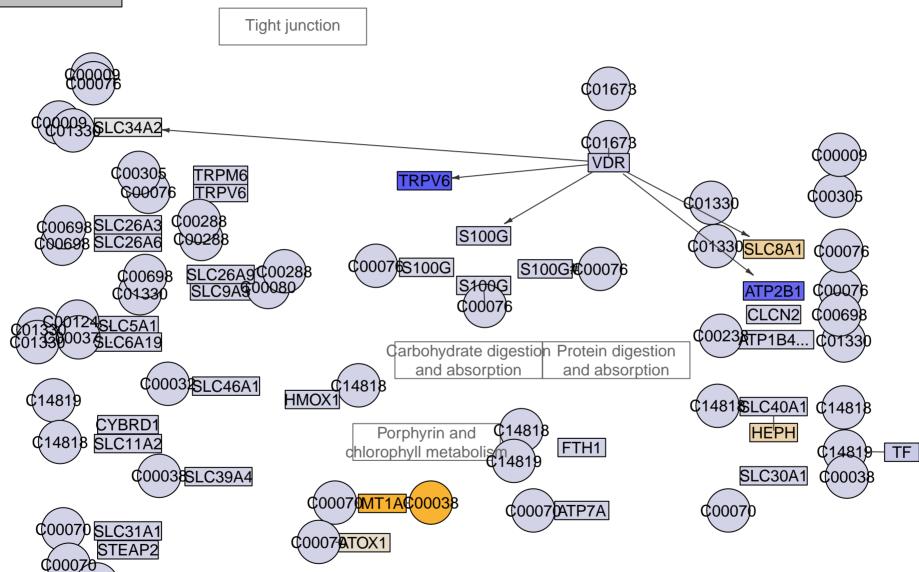


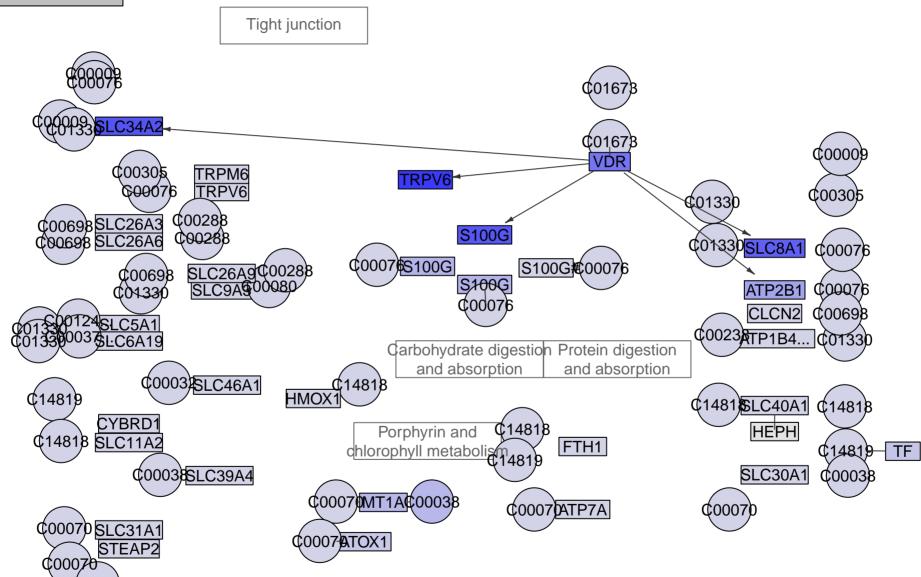
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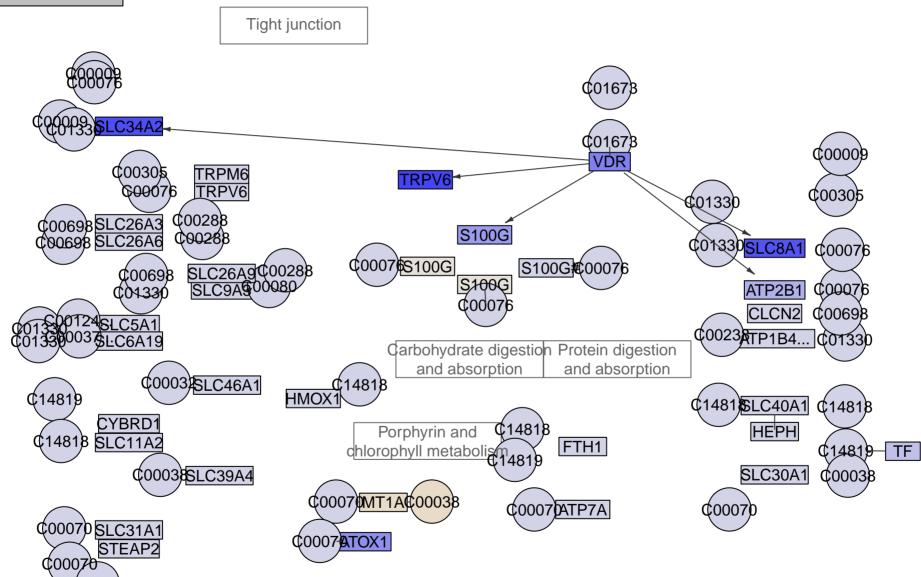
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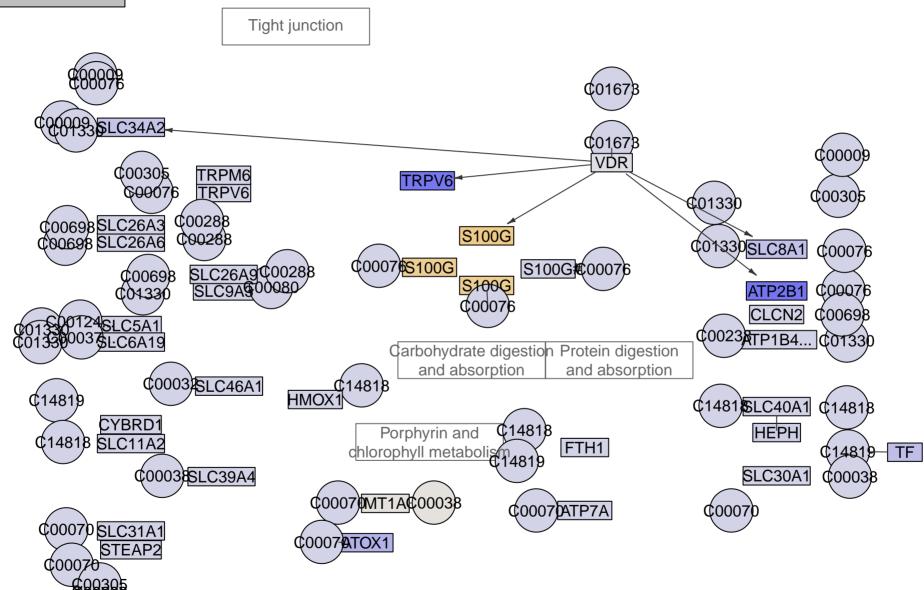


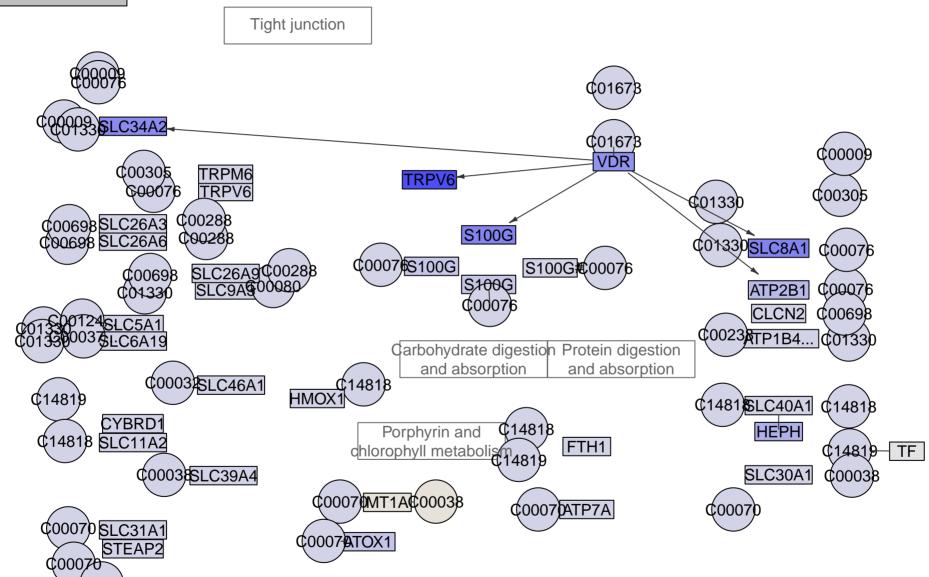
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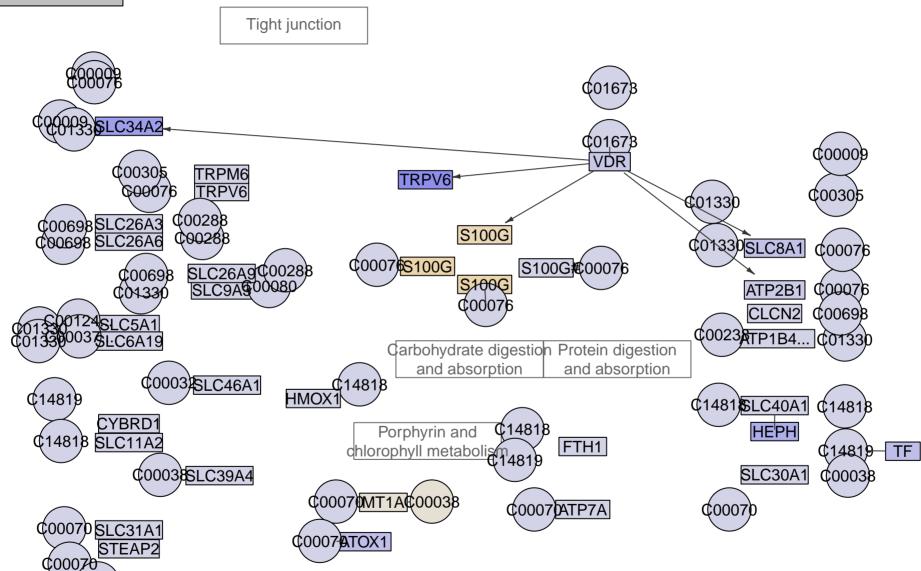


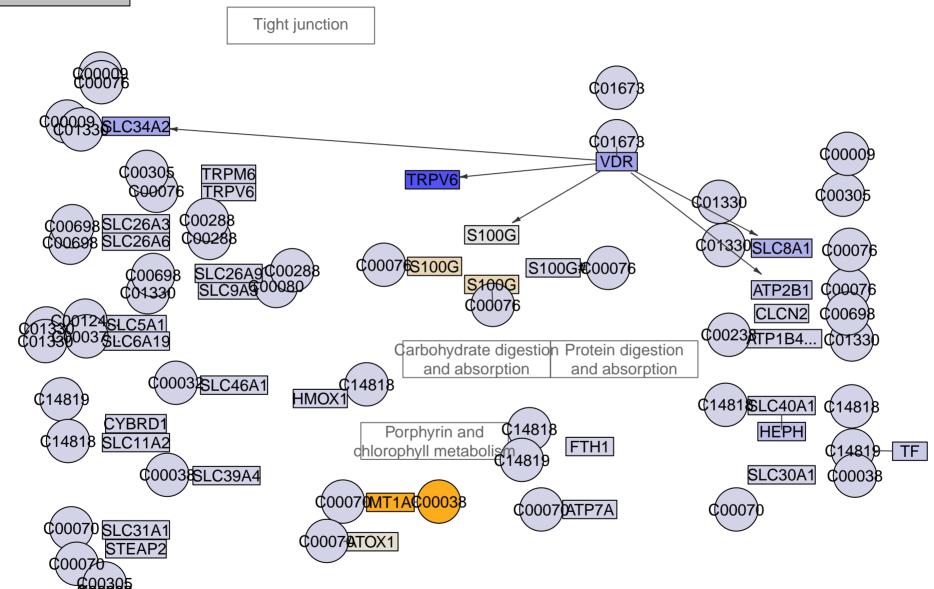


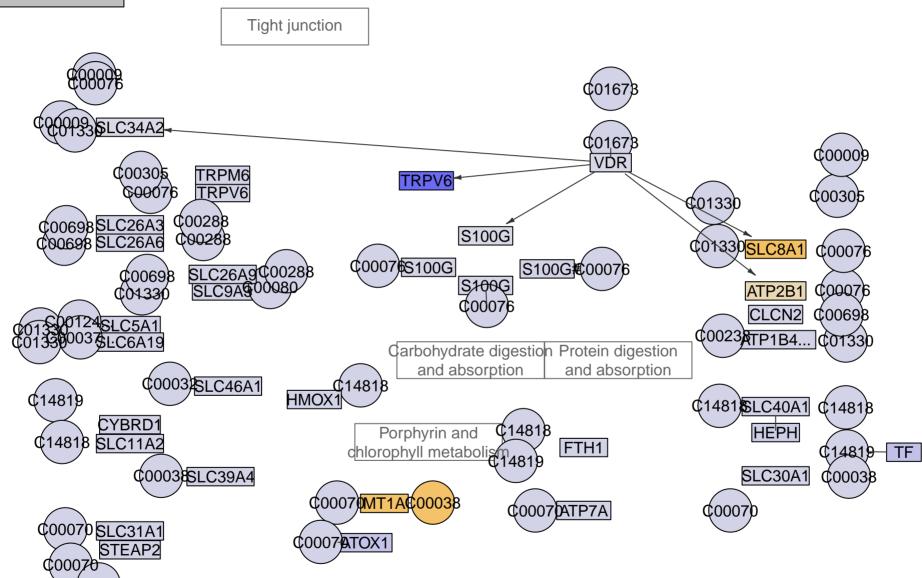


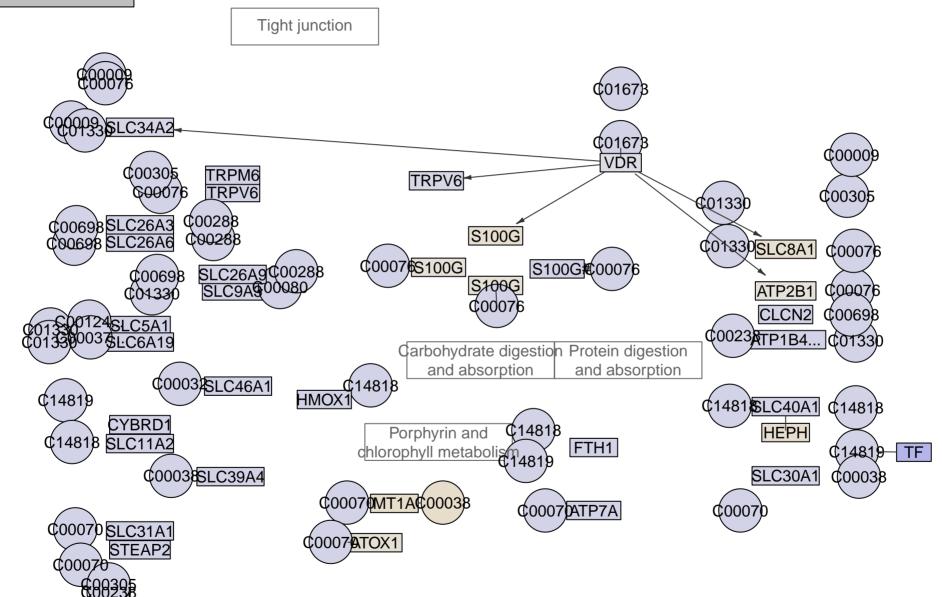




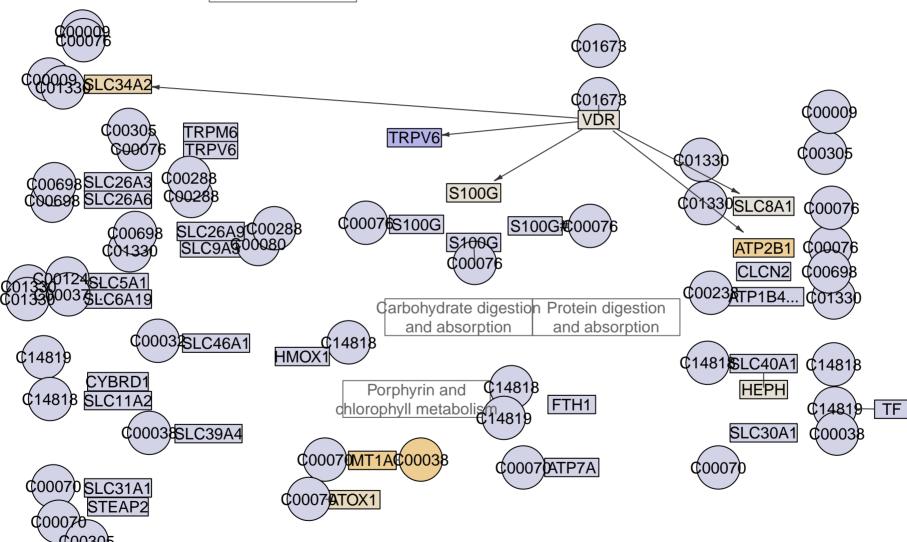


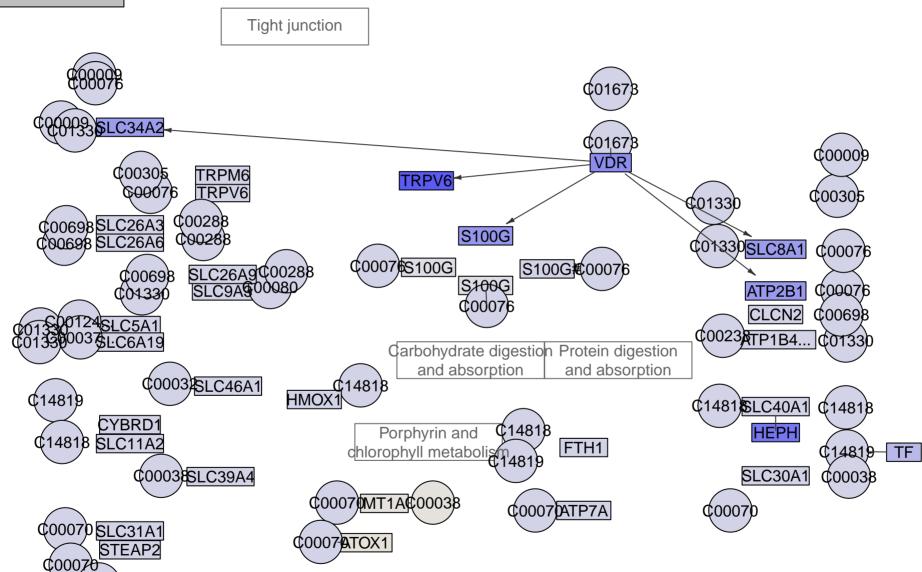


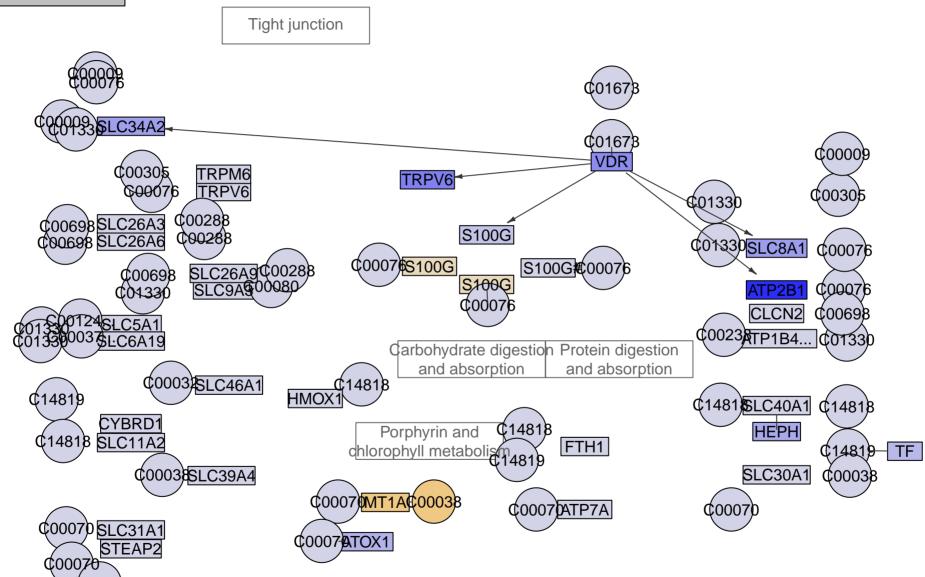


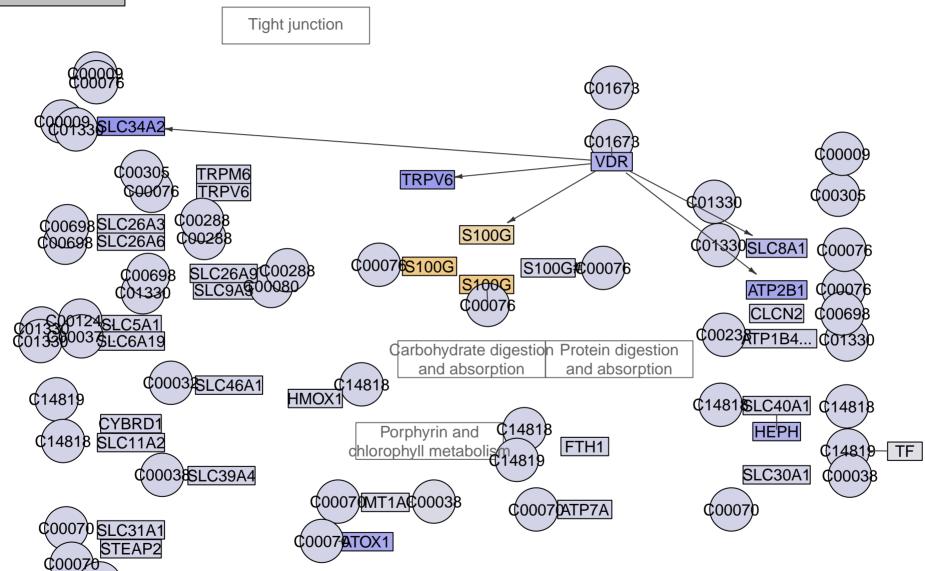


Tight junction

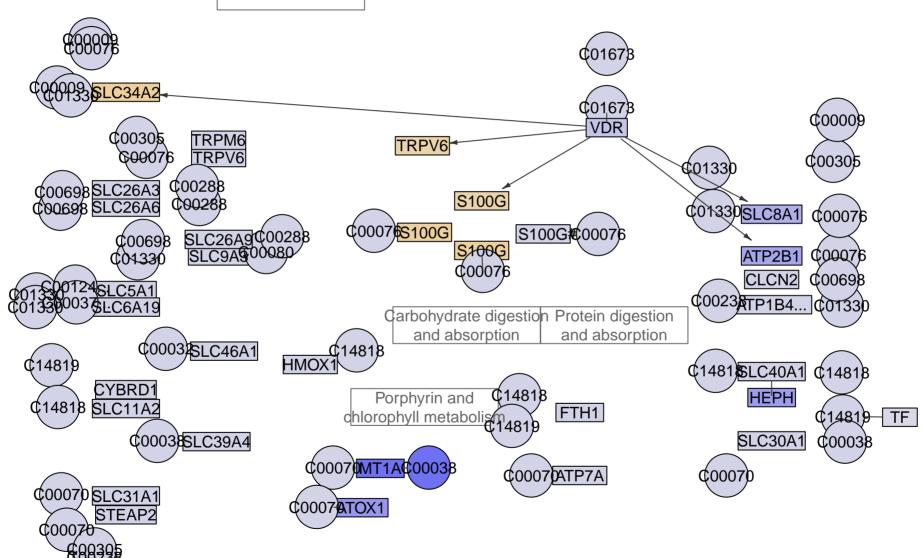


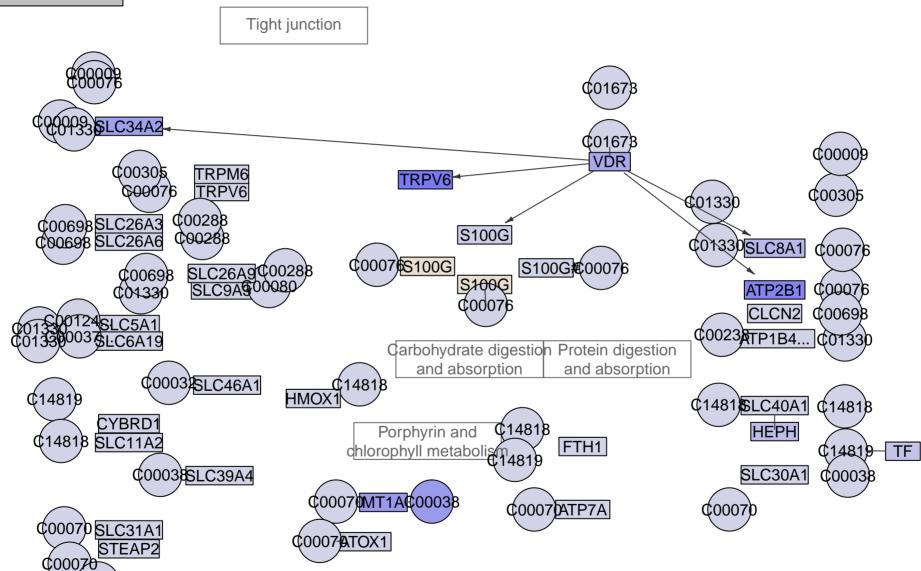


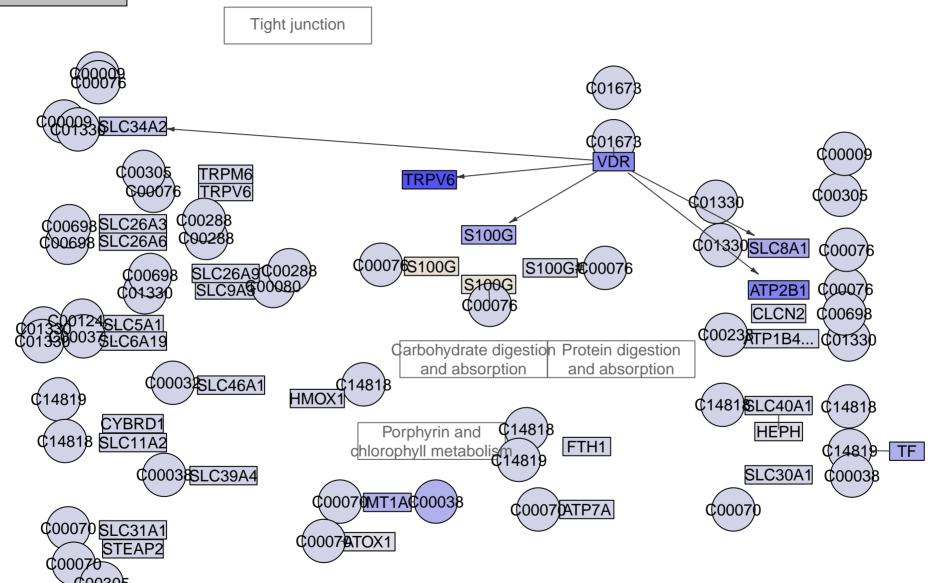


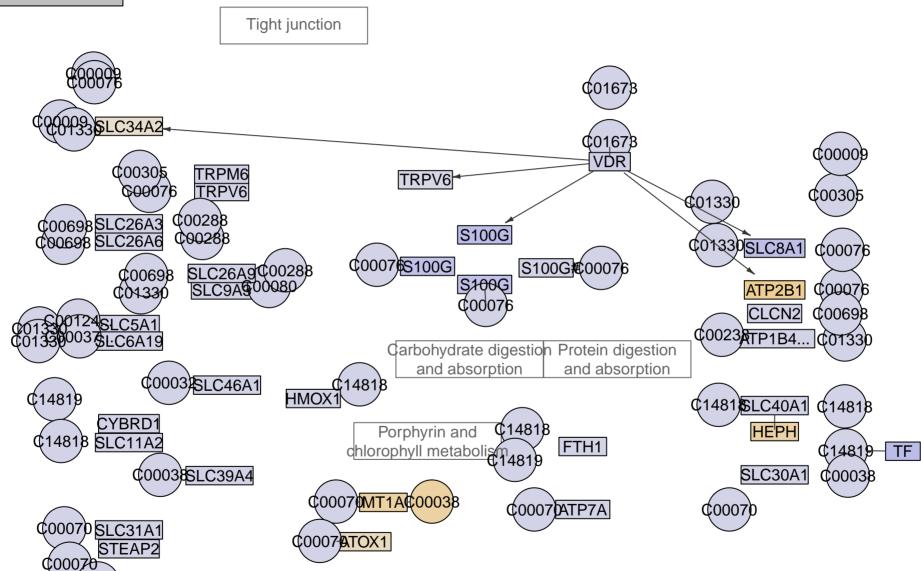


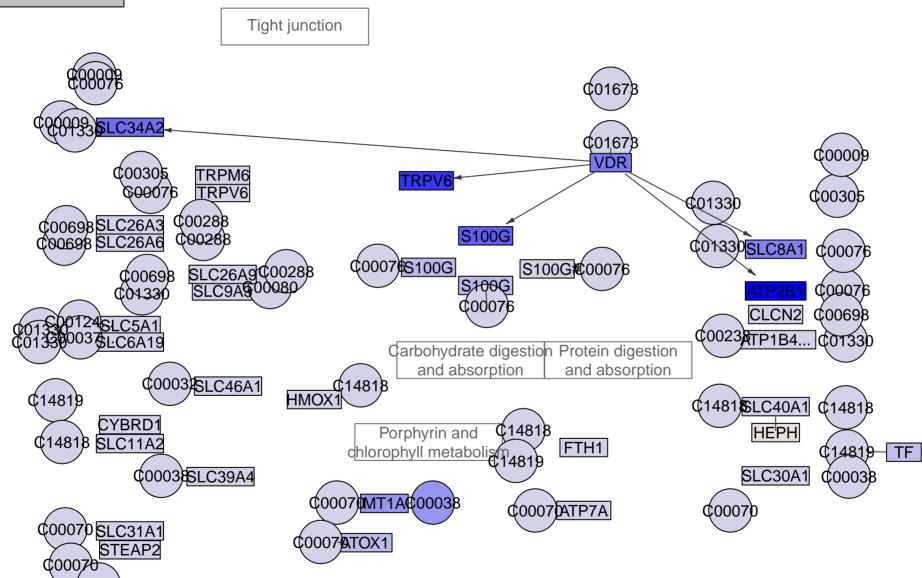


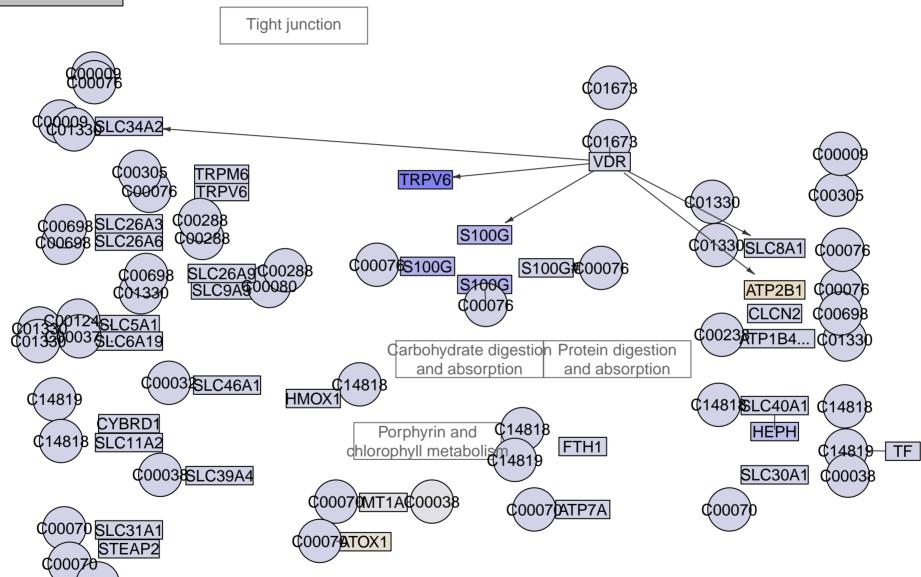


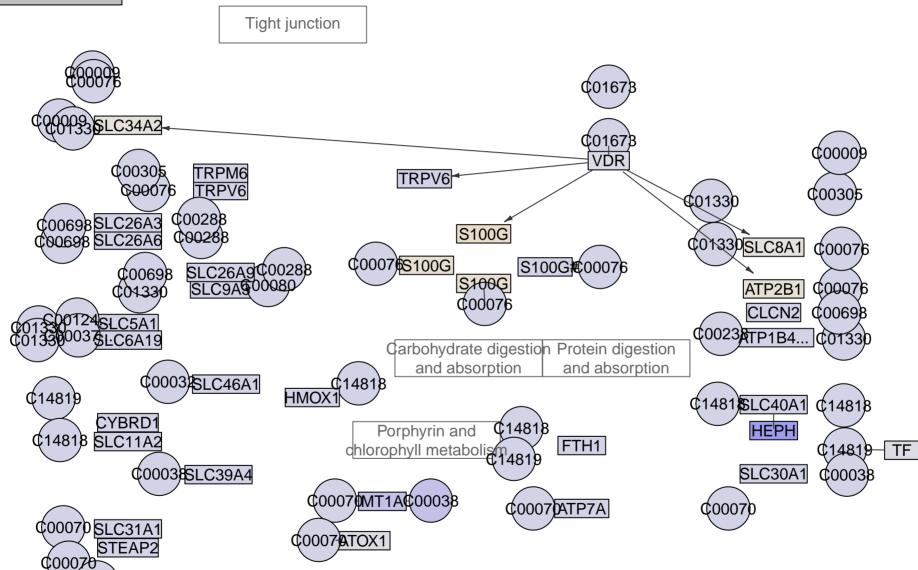








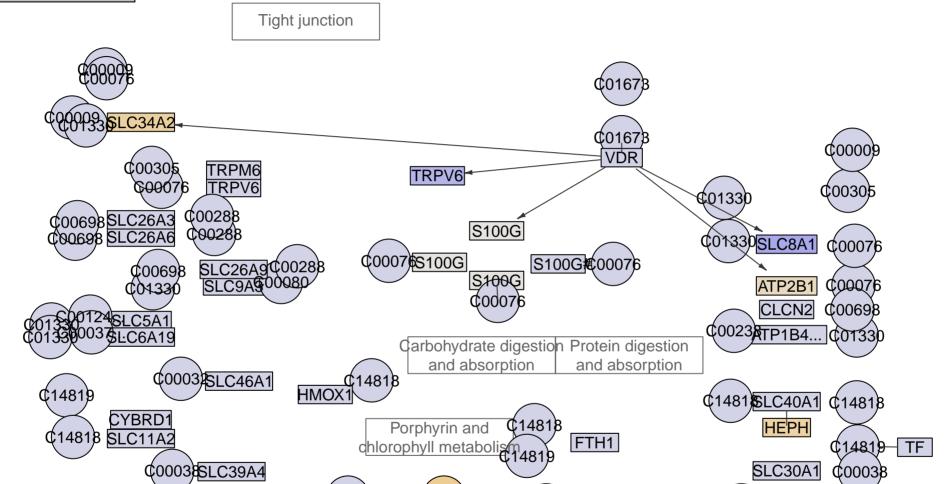




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STEAP2

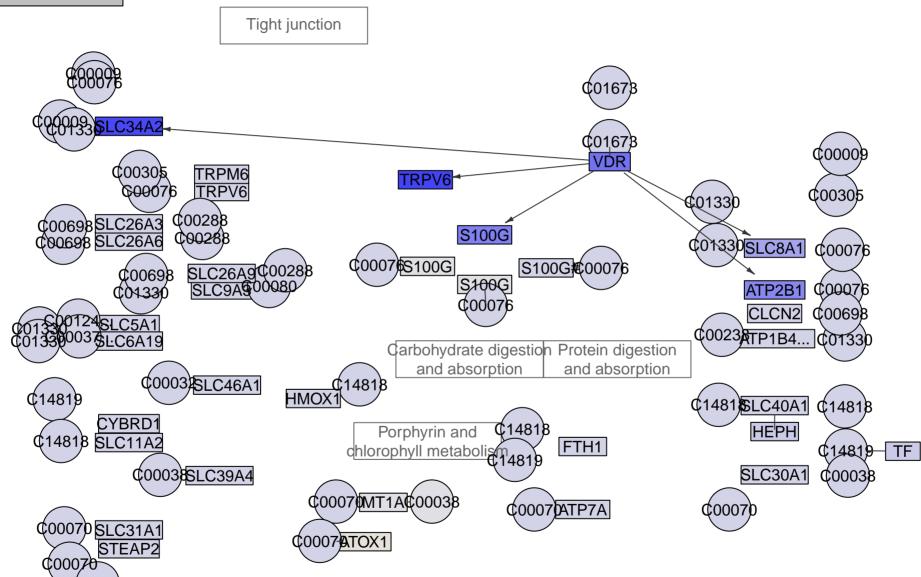


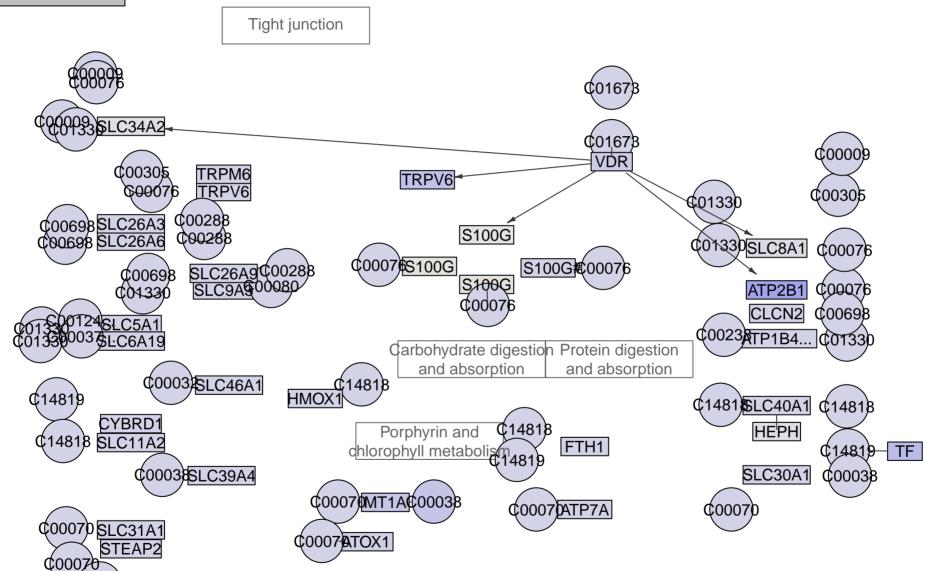
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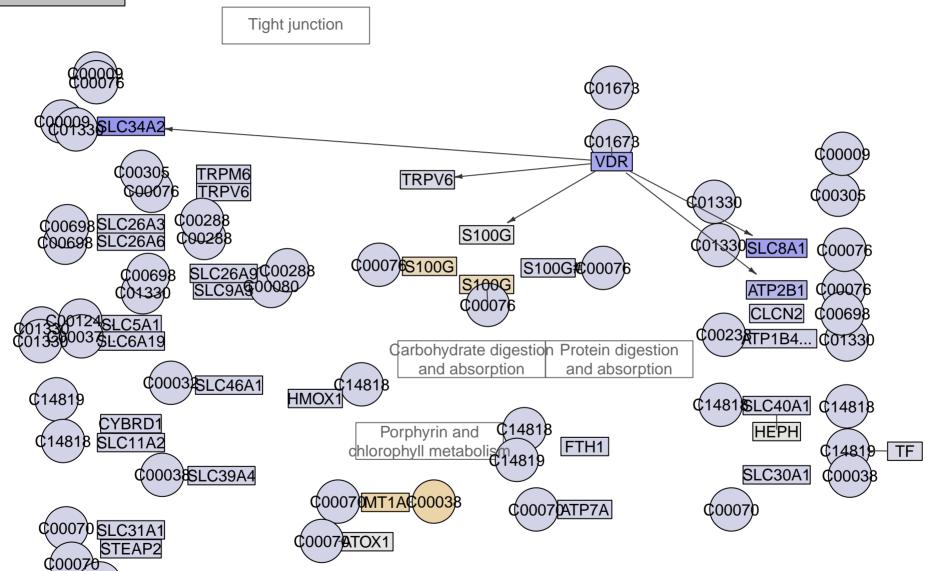
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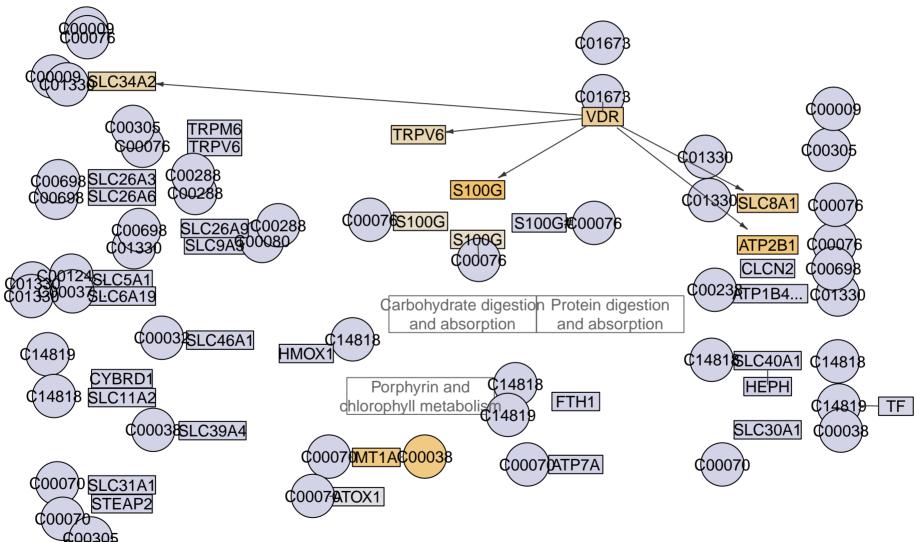
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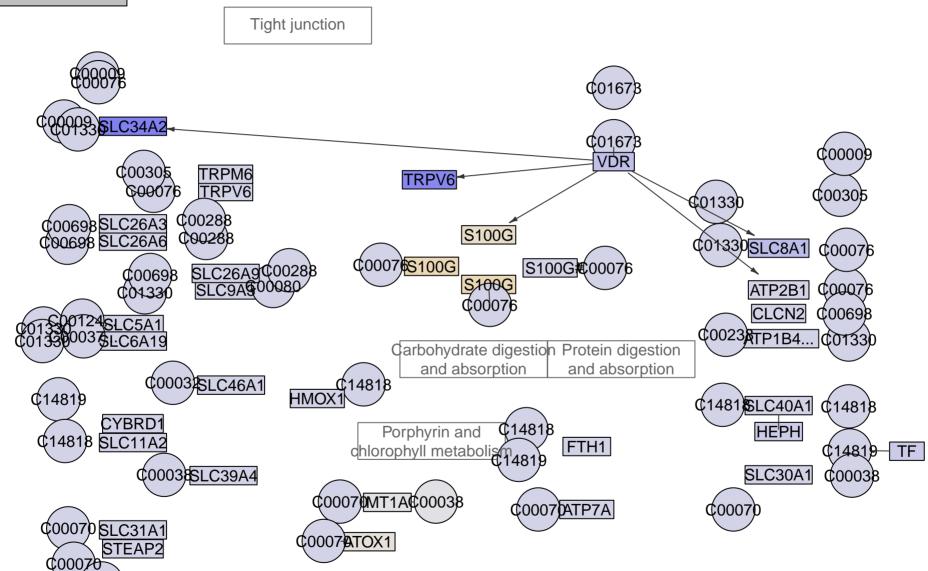


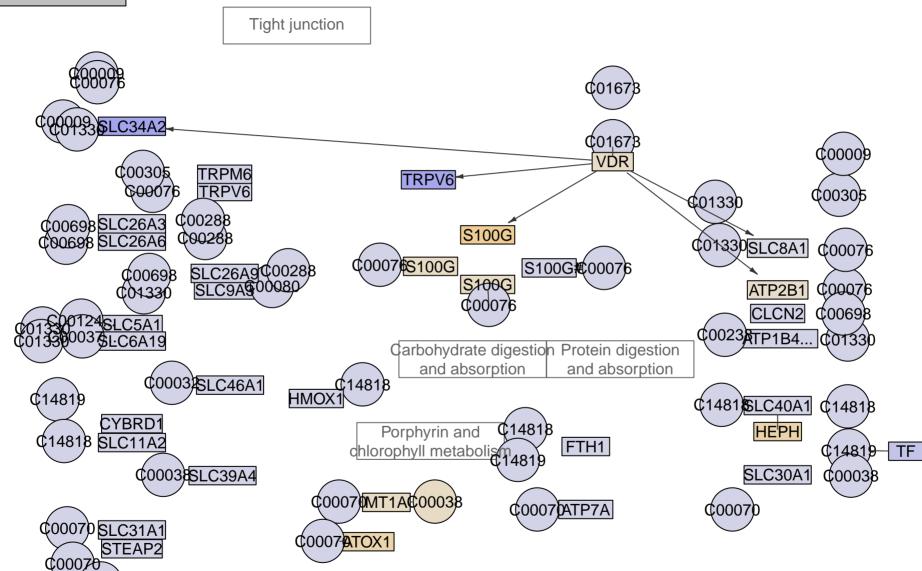


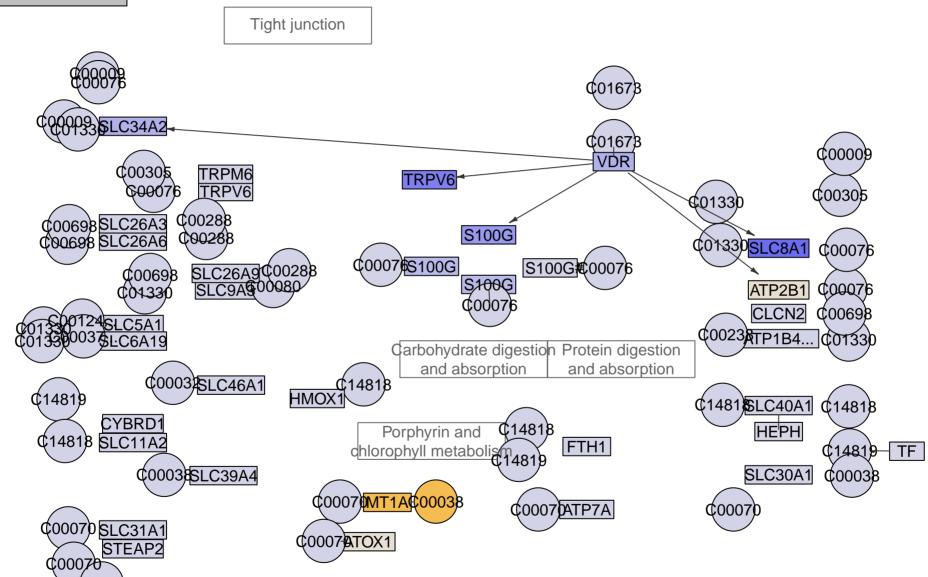


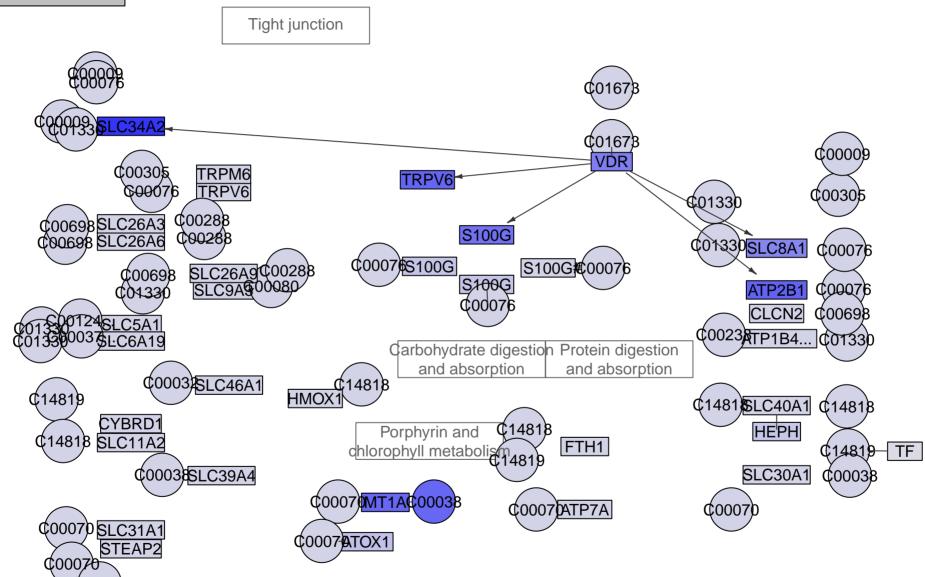
Tight junction

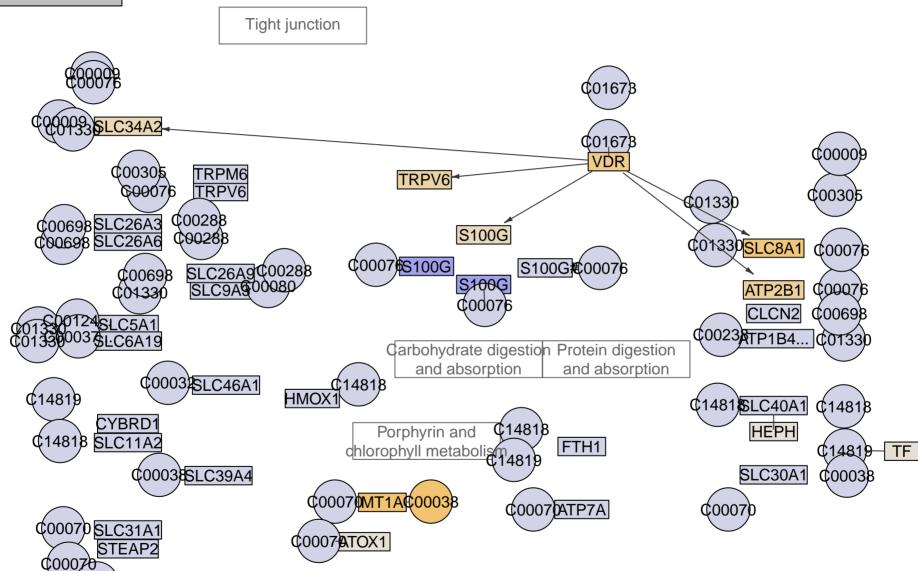


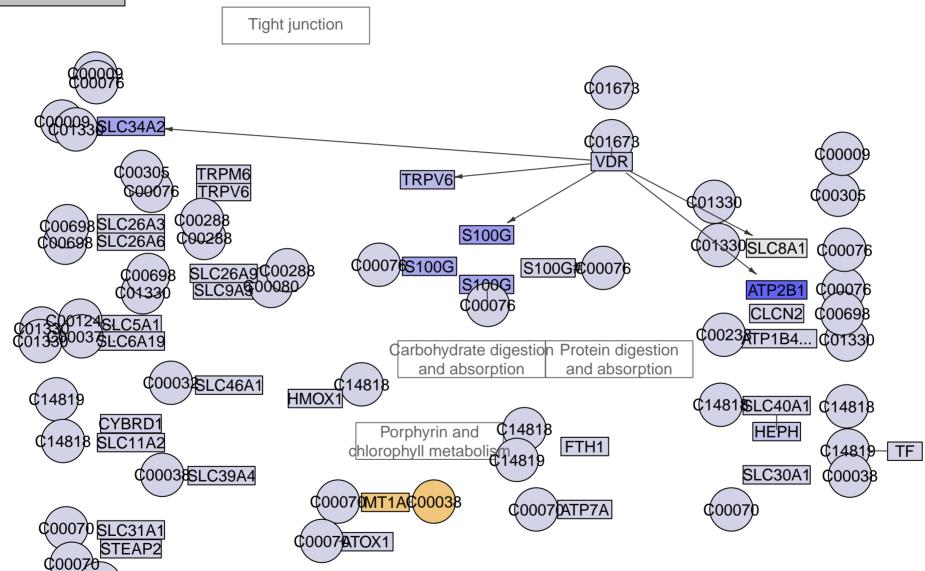


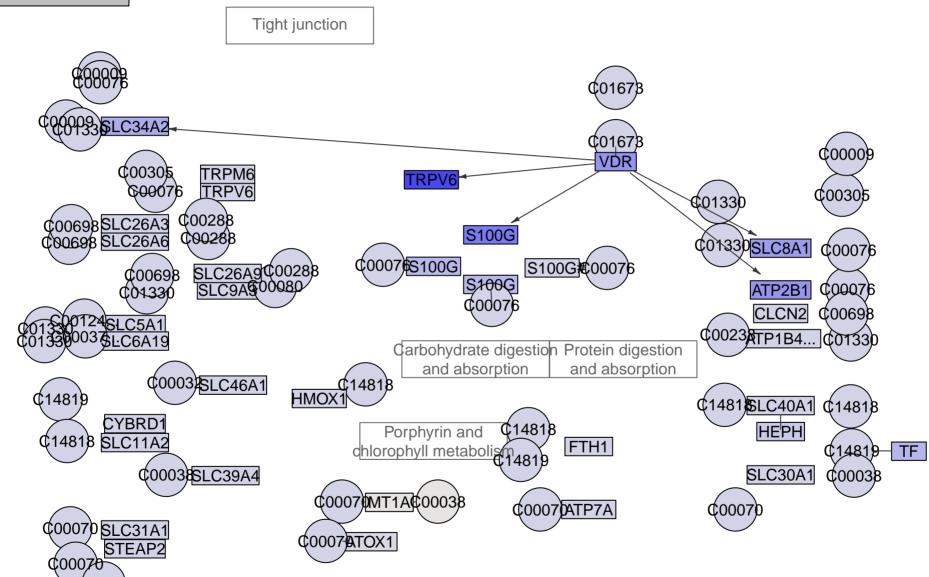


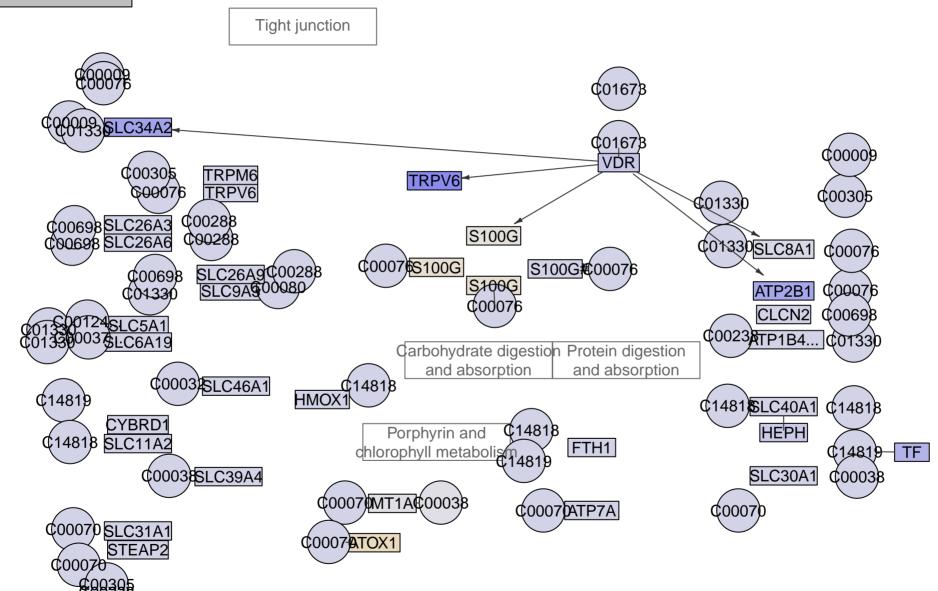


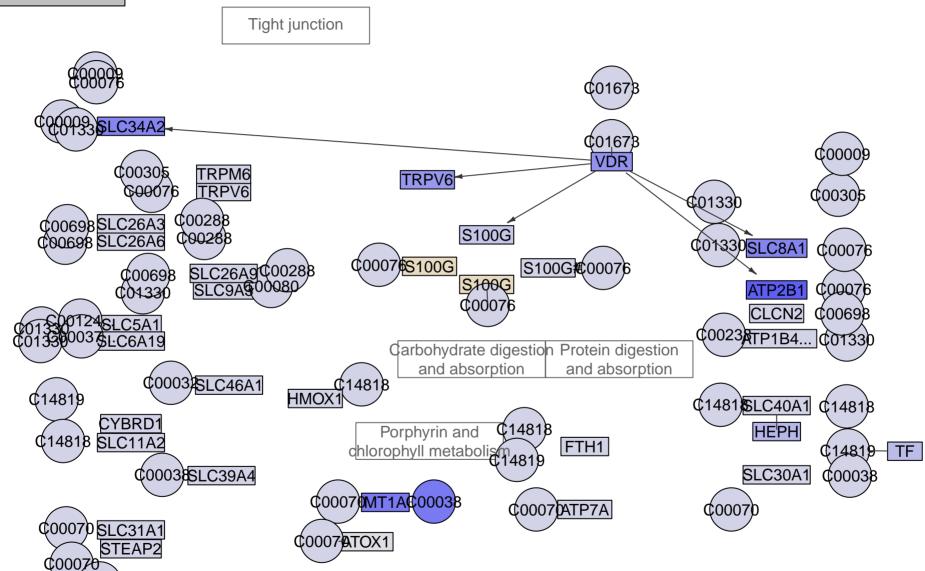


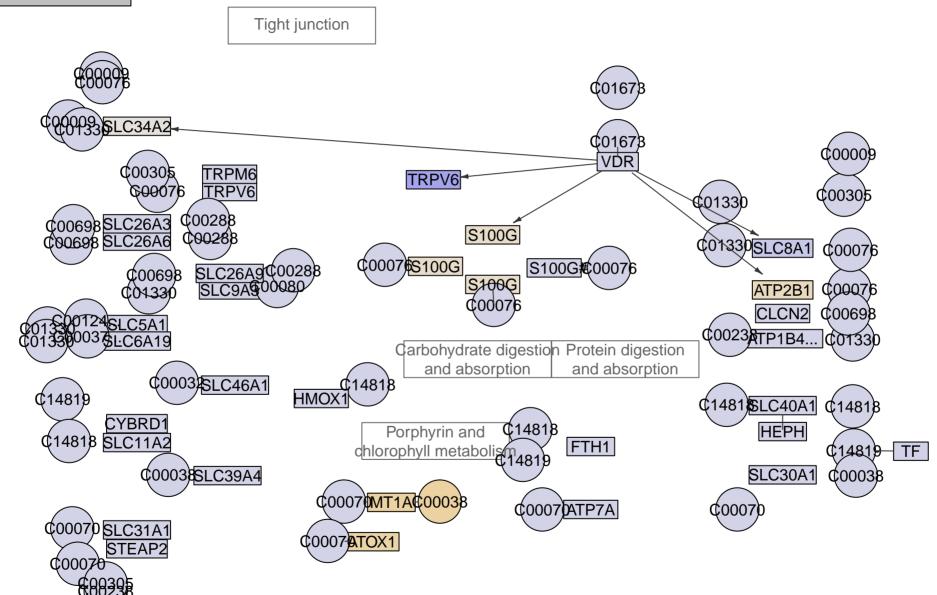


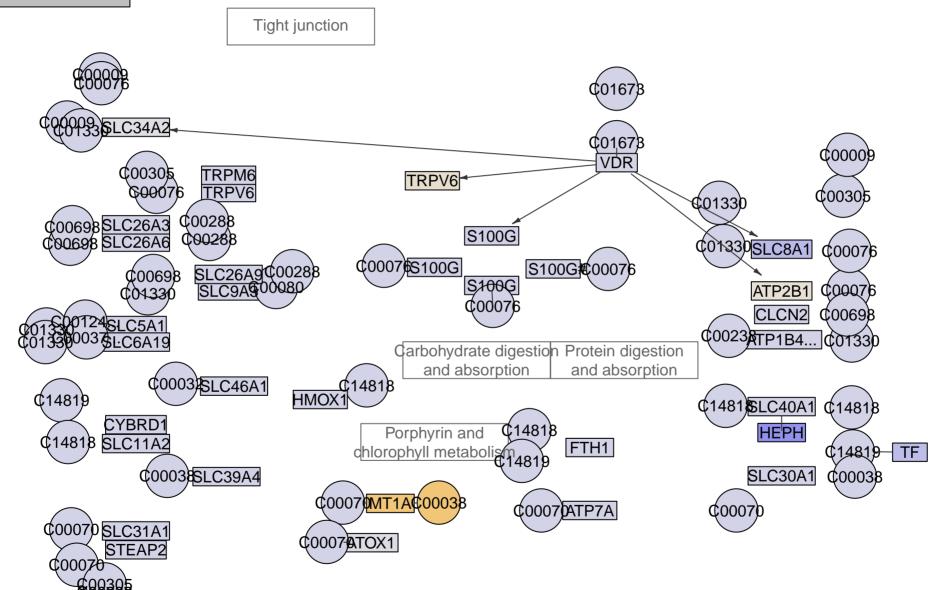


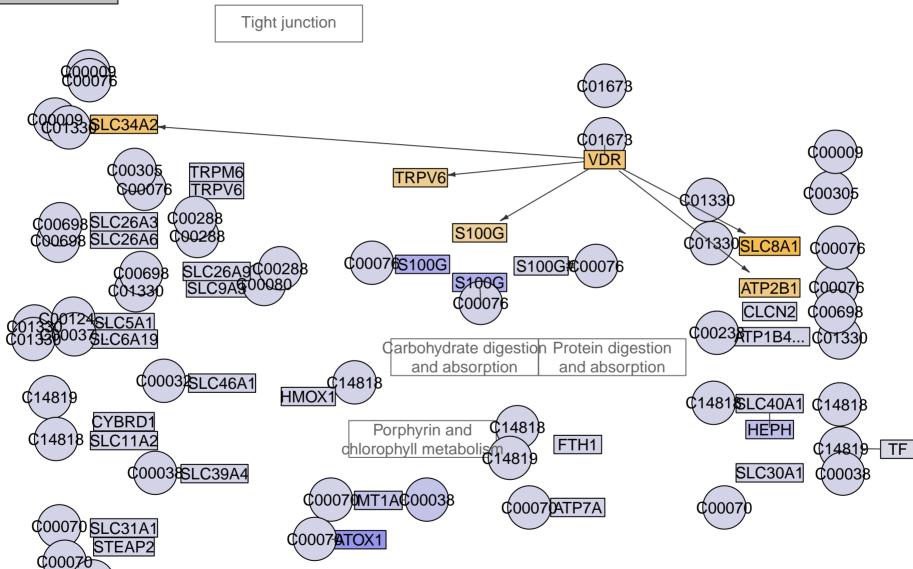


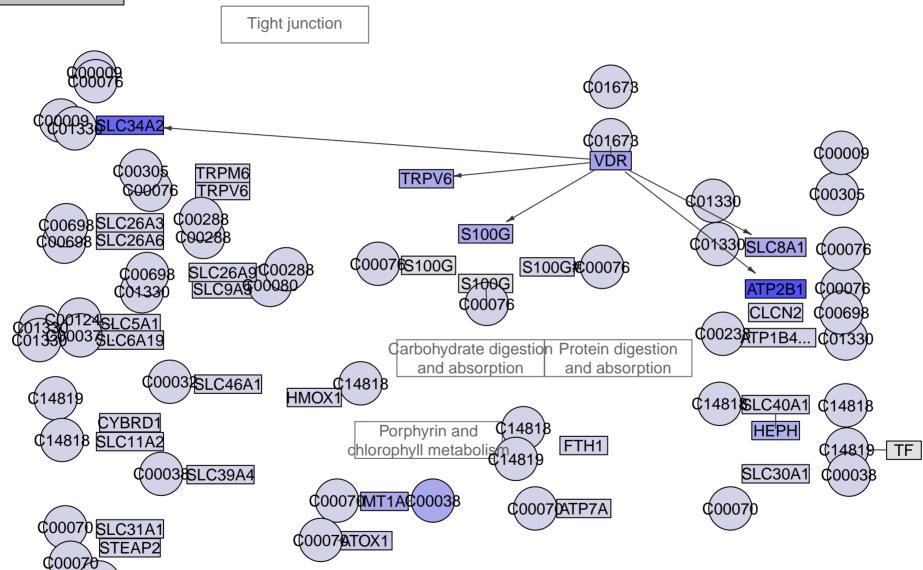


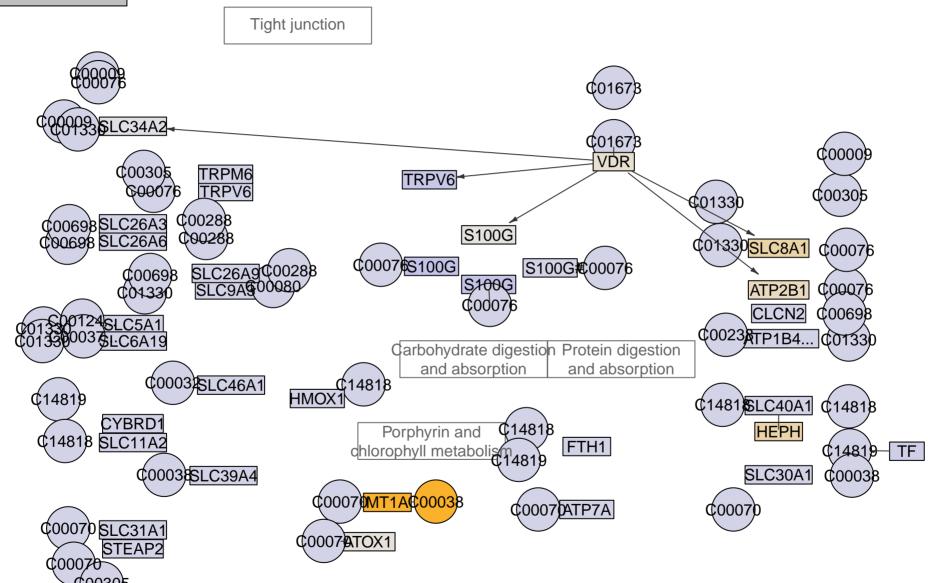




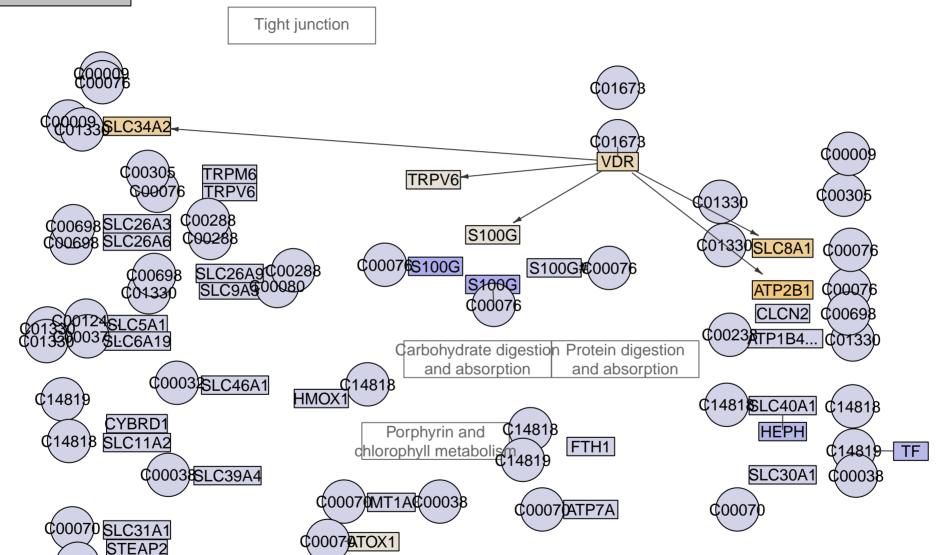


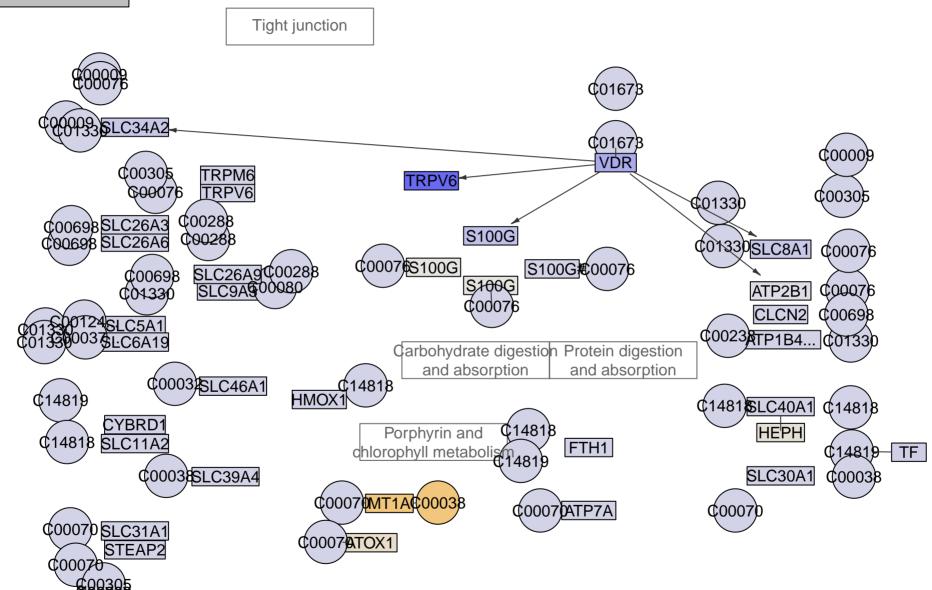


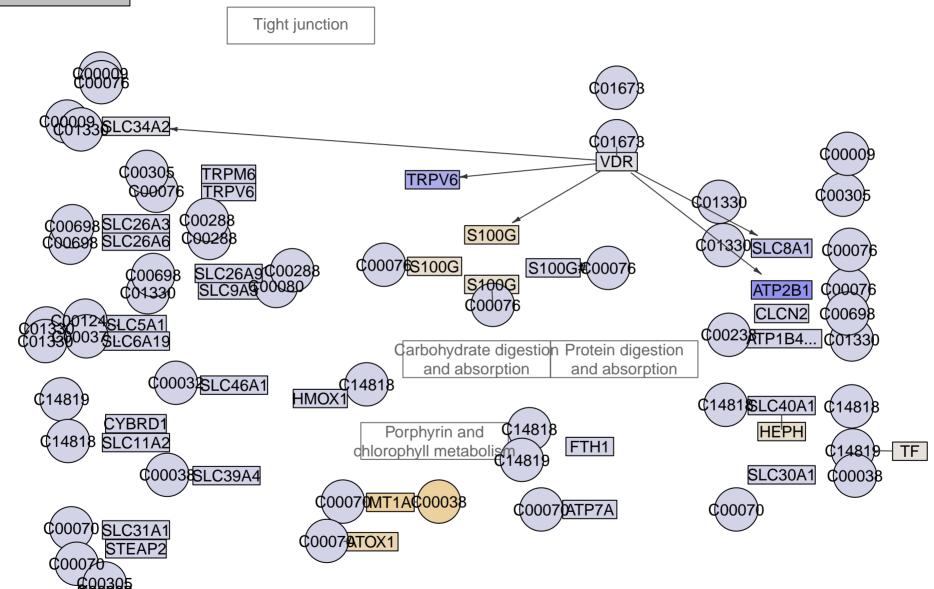




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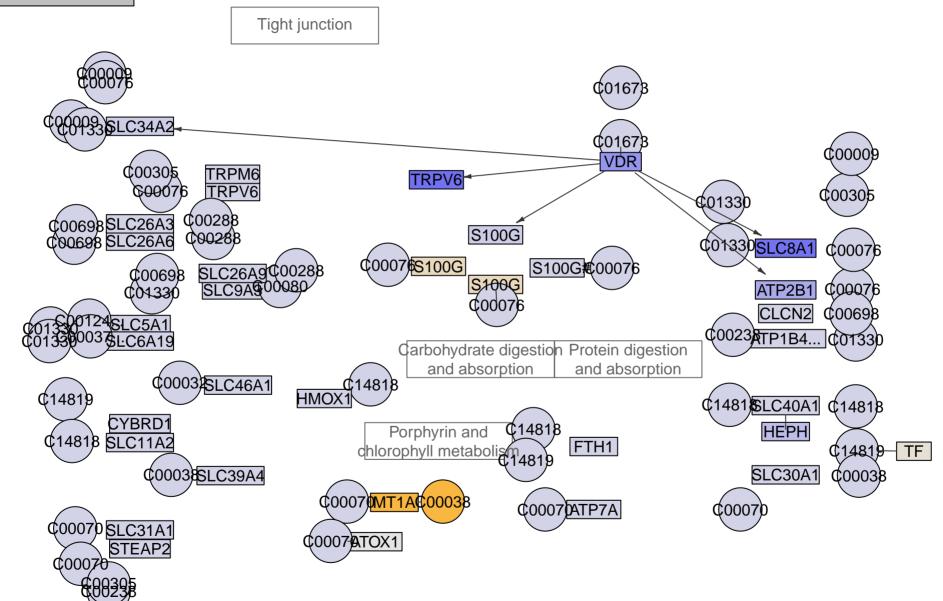




Mineral absorption MPI-129 Tight junction **¢**0167 C0167 ¢0000 VDR C0030 TRPM6 TRPV6 TRPV6 600076 ¢00305 ¢01330 ¢00288 C00698 SLC26A3 C00698 SLC26A6 S100G ¢90288 ¢01330<mark>SLC8A1</mark> ¢00076 ¢0007\$\$100G SLC26A9(0028) SLC9A(00080 S100G#00076 ¢00698 S100G ATP2B1 C00076 **¢**04330 C00076 CLCN2 C00698 00124SLC5A1 350003 \$LC6A19 ¢0023 C01330 Carbohydrate digestion Protein digestion and absorption and absorption C0003 SLC46A1 ¢14819 HMOX C1481 SLC40A1 C14818 CYBRD<sup>2</sup> C14818 Porphyrin and HE<sup>PH</sup> chlorophyll metabolis (1481) C14818 SLC11A2 FTH1 ¢14819 TF C0003 SLC39A4 SLC30A1 ¢00038 C00070MT1AC00038 C0007)ATP7A (0007) ¢00070 SLC31A1 **¢**0007**€**TOX1

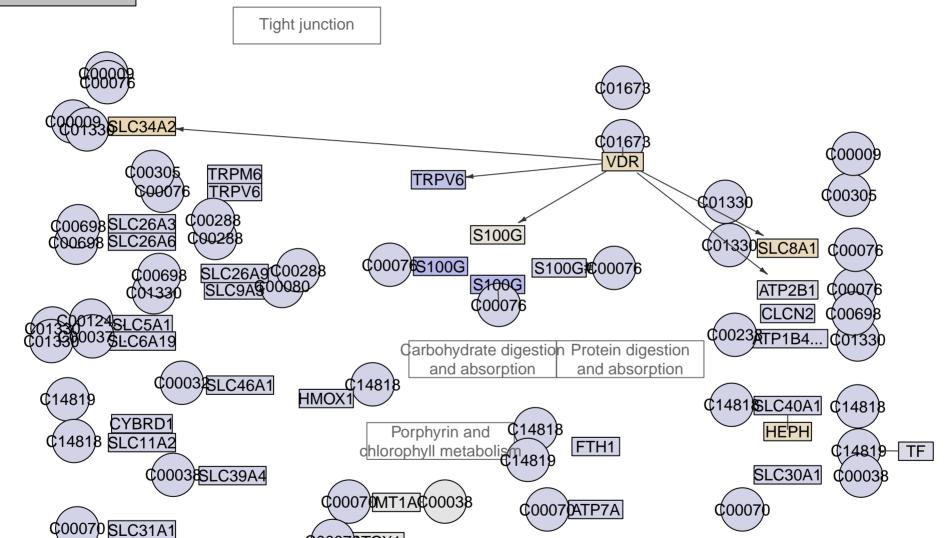
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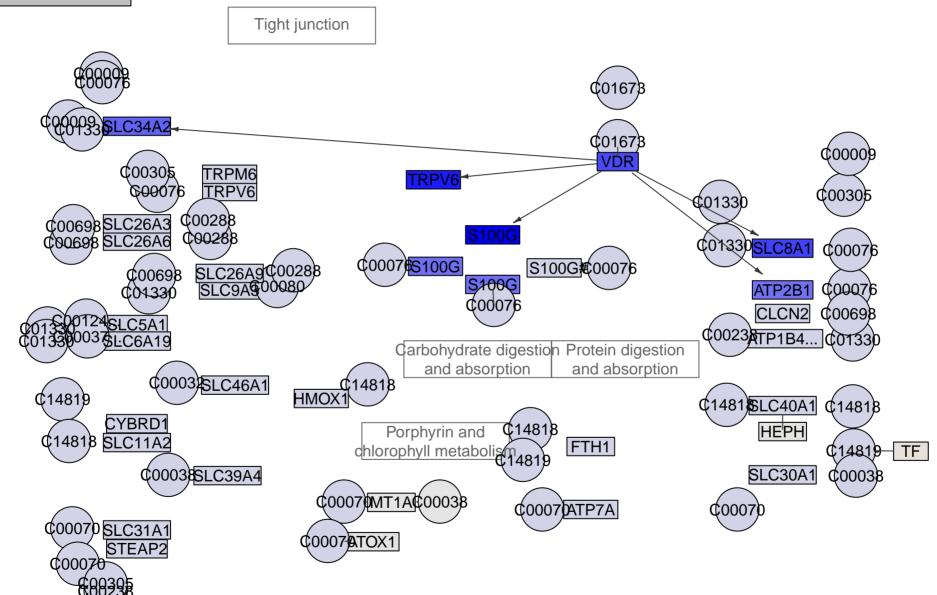


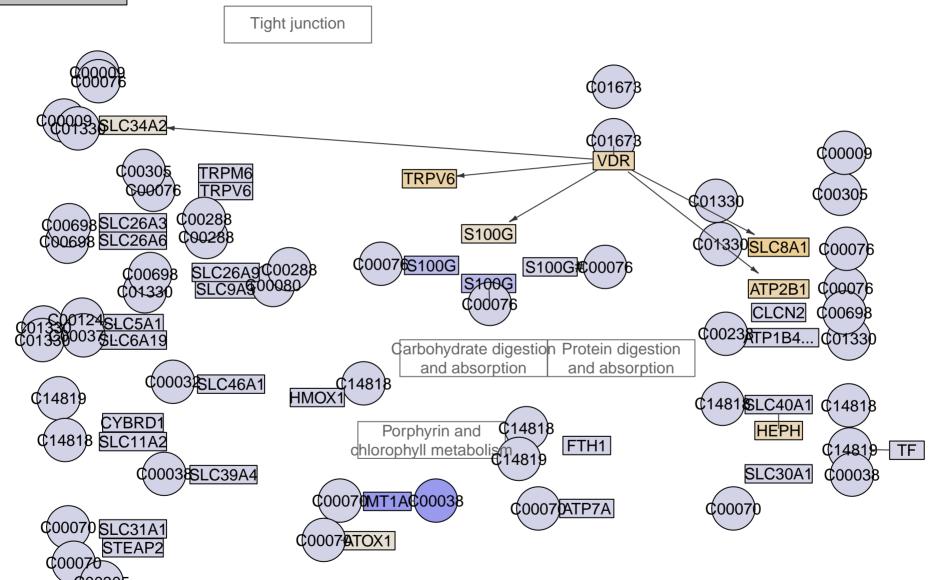
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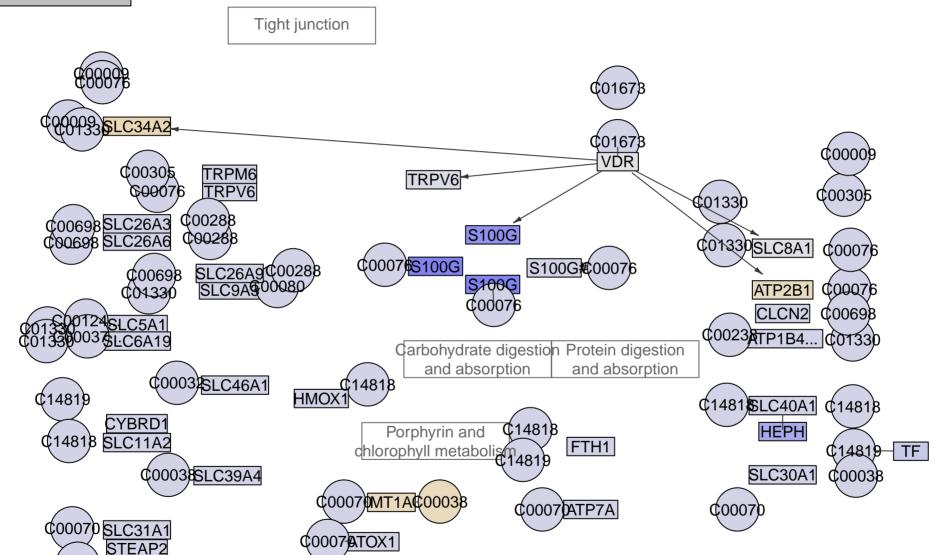


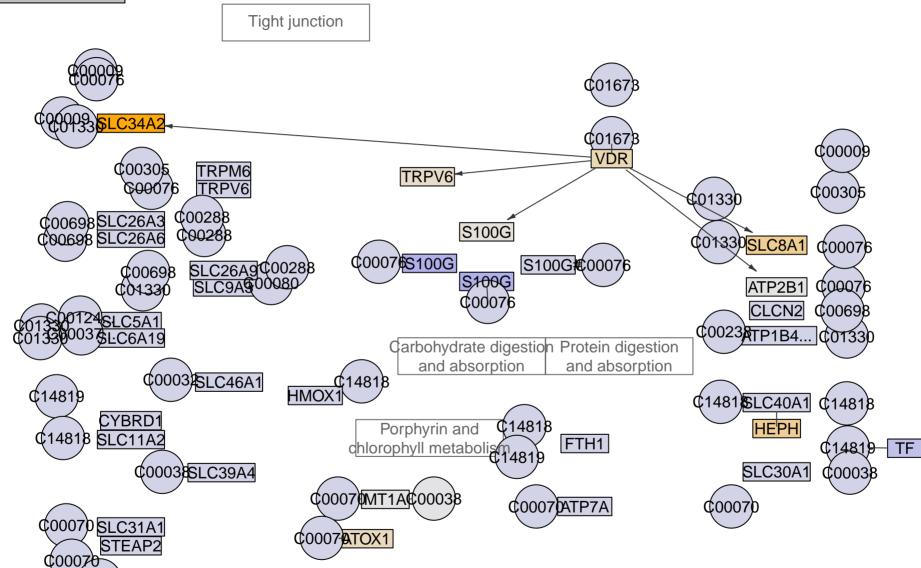
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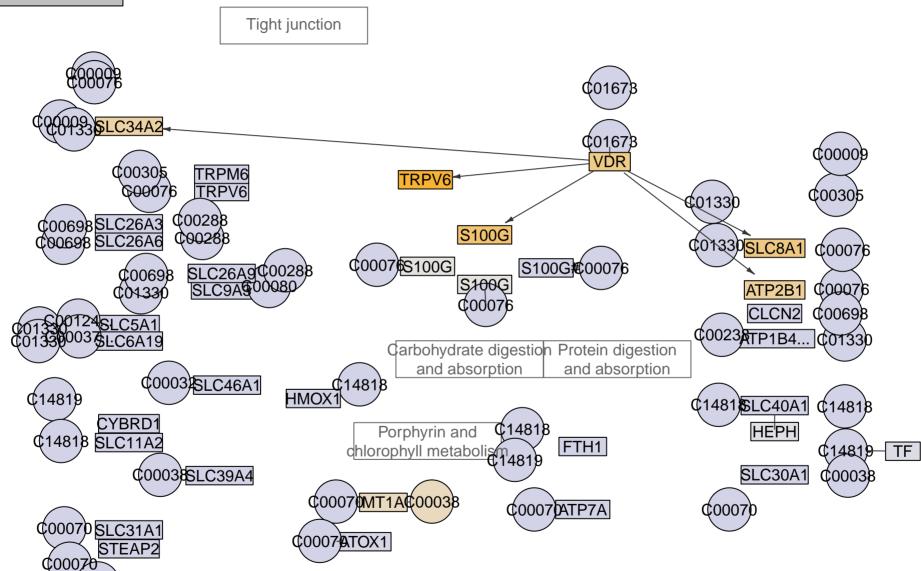


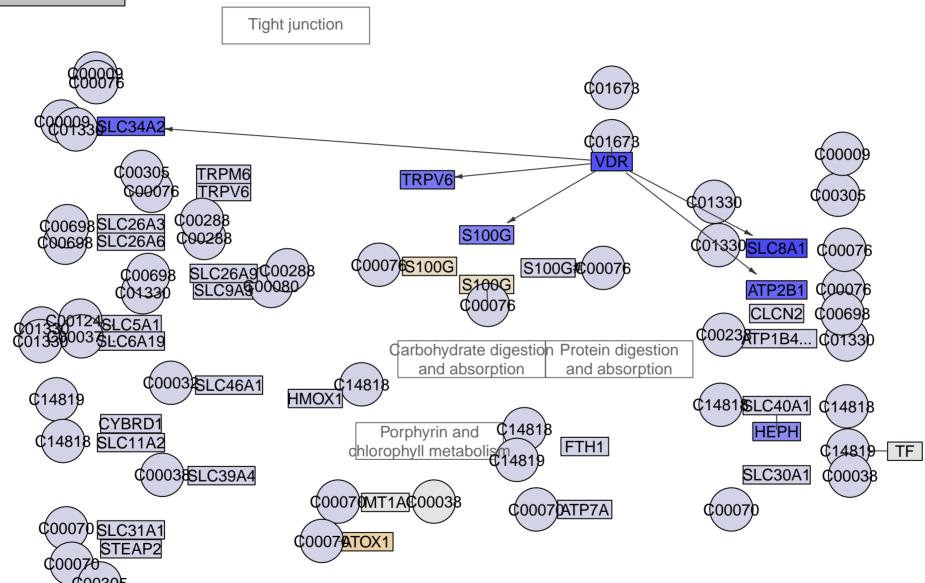


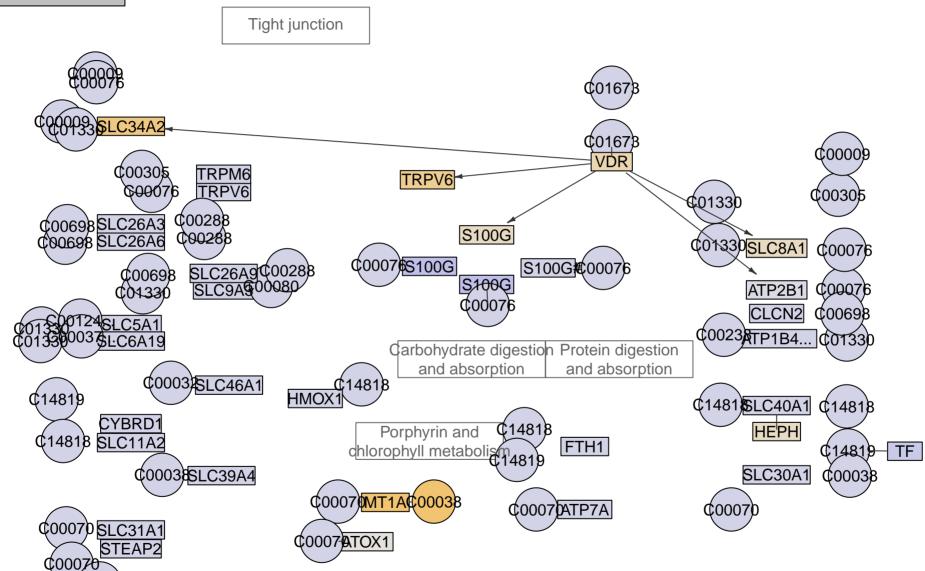
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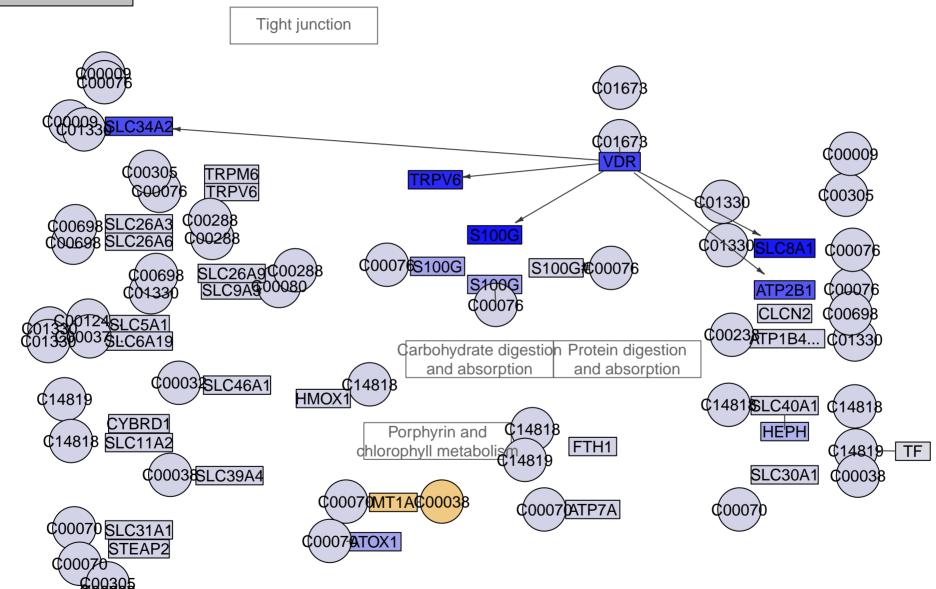


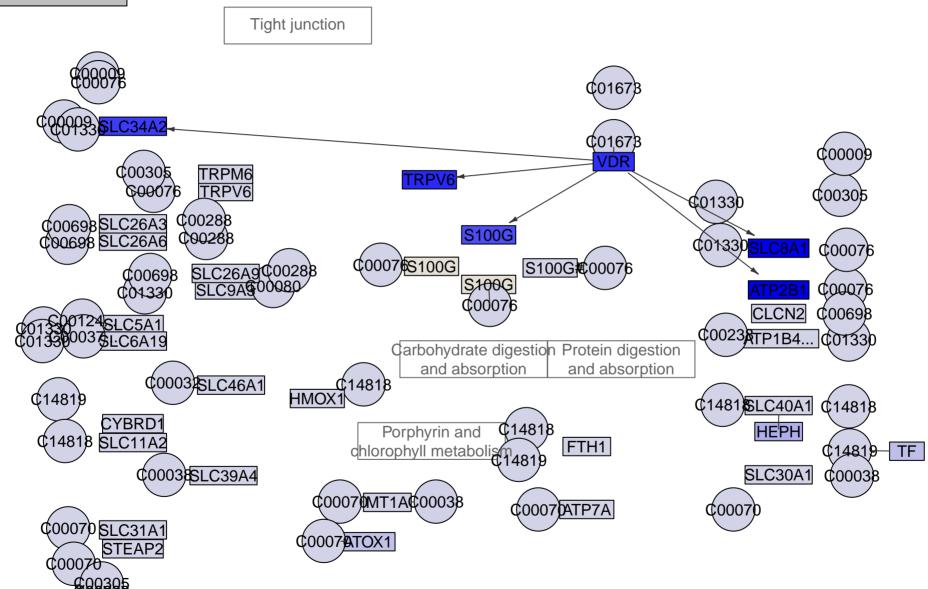


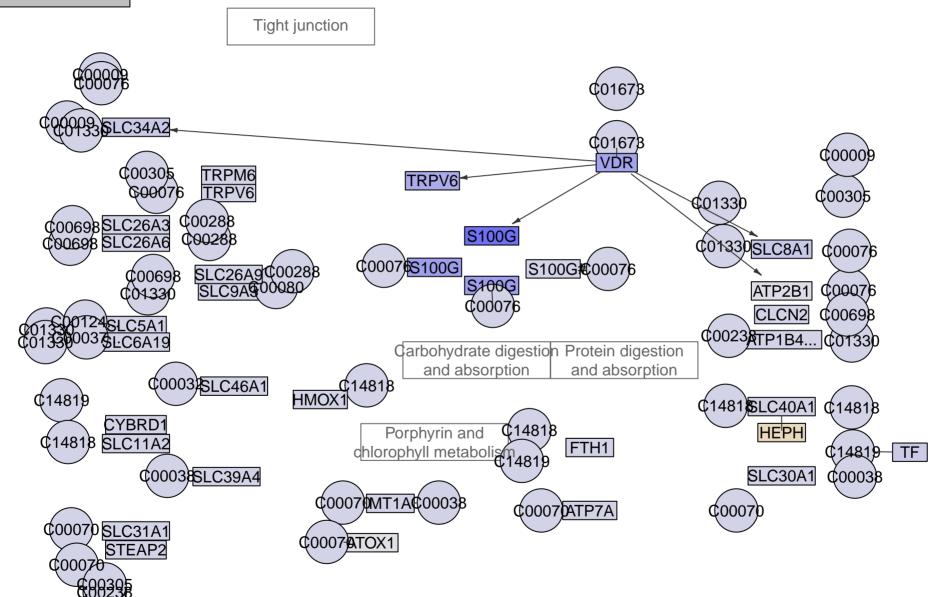




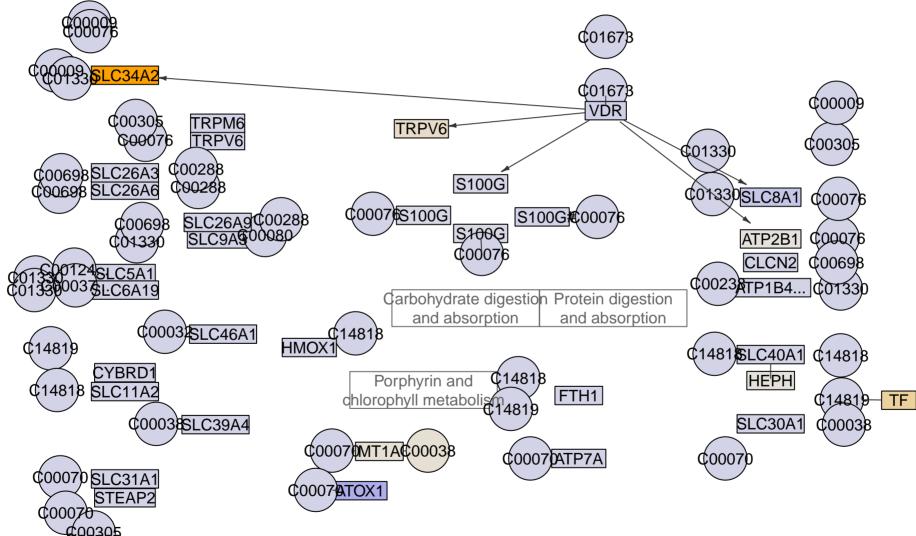


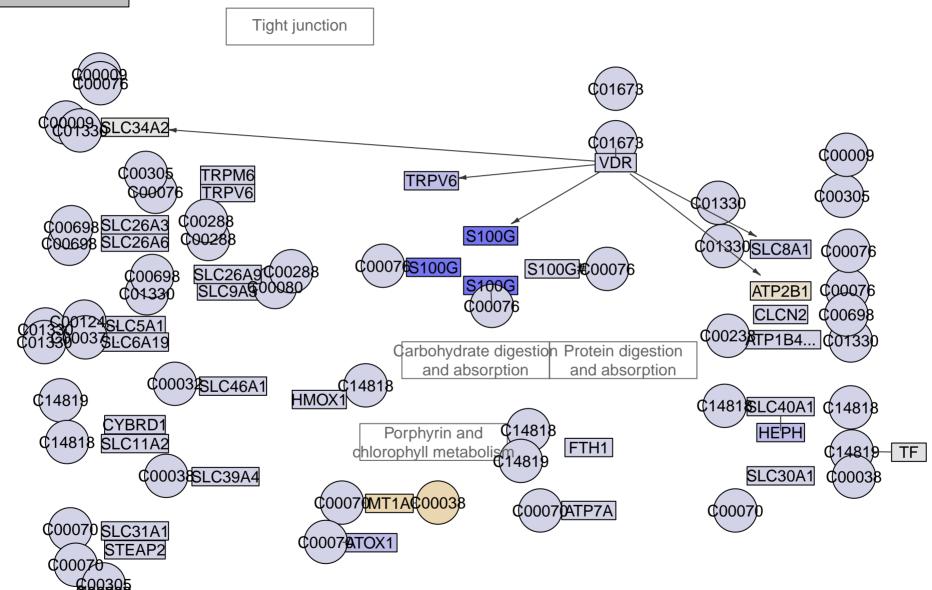


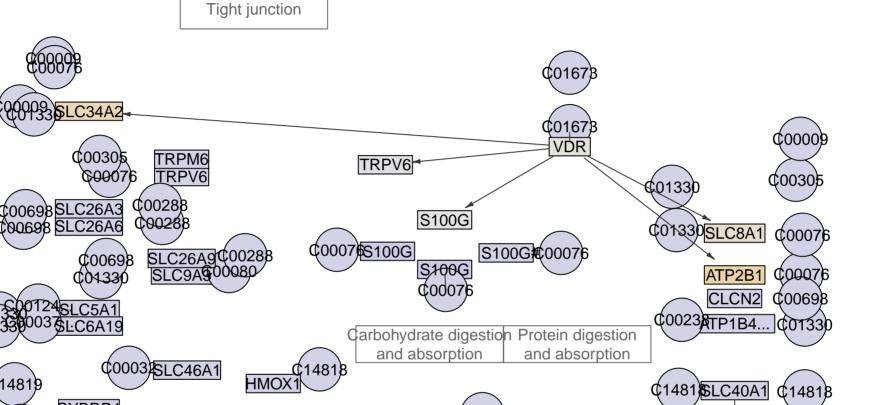


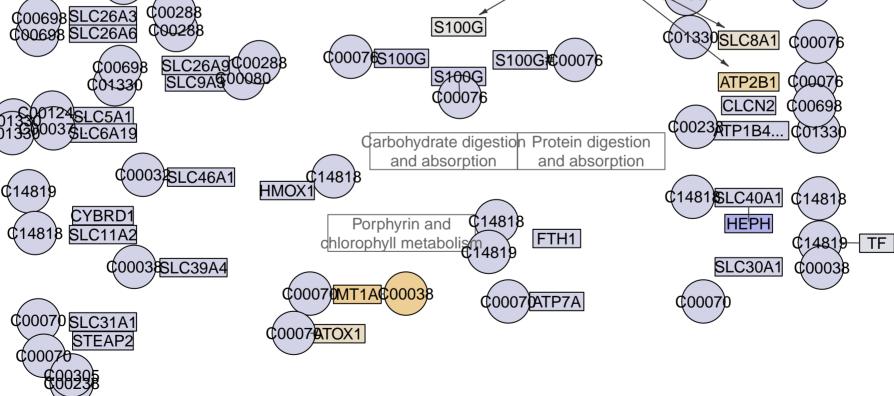


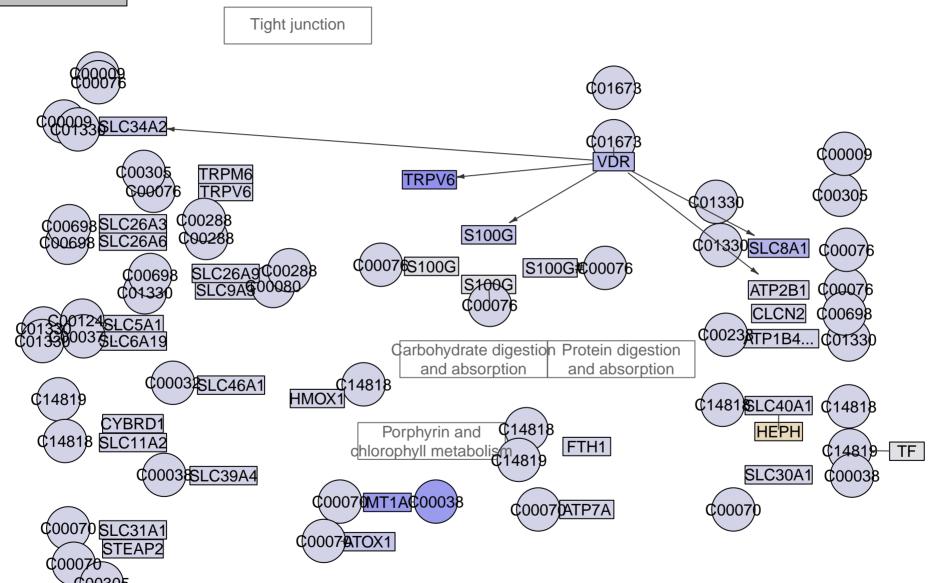


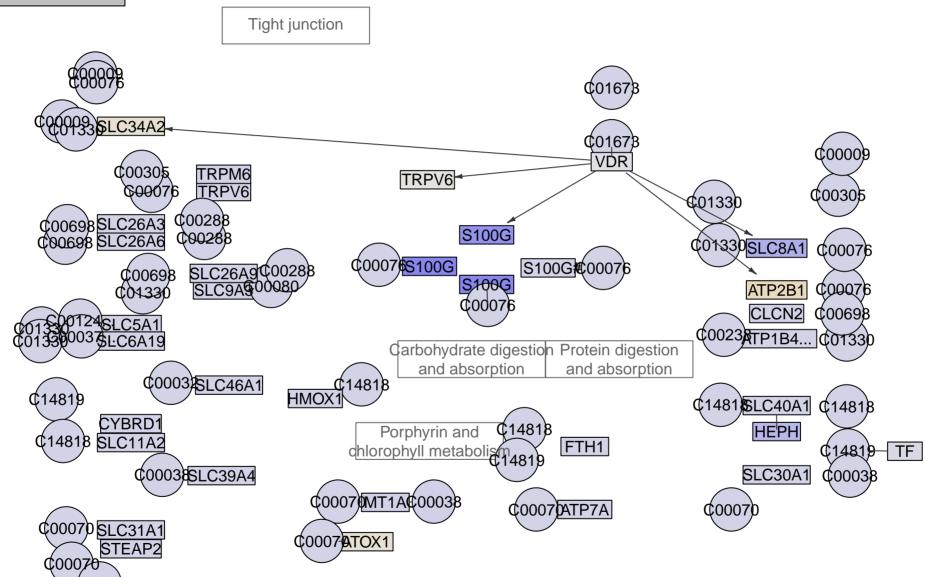


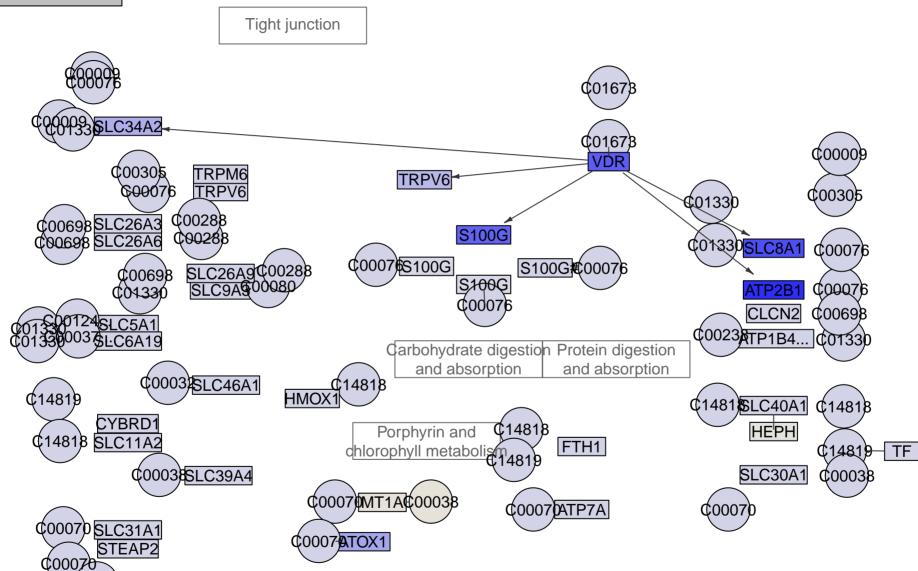


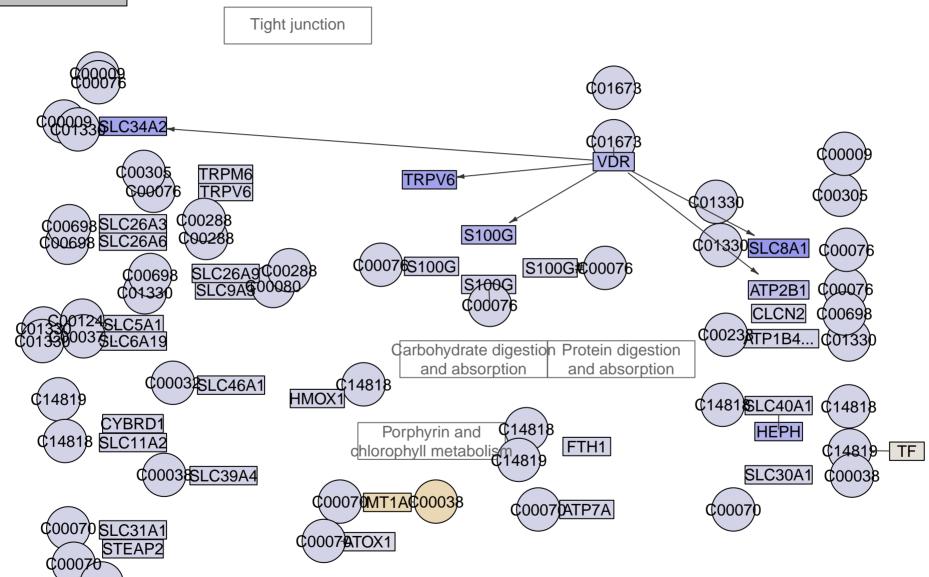


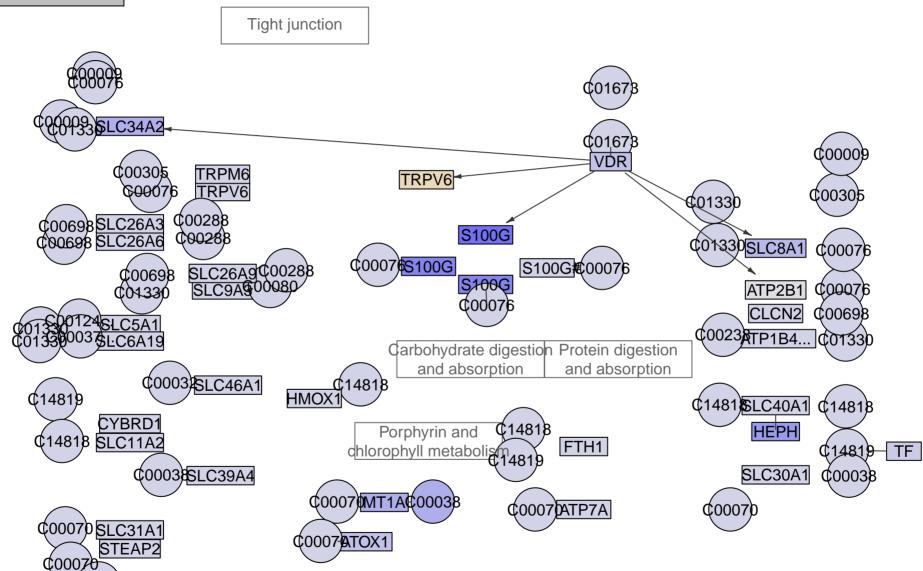




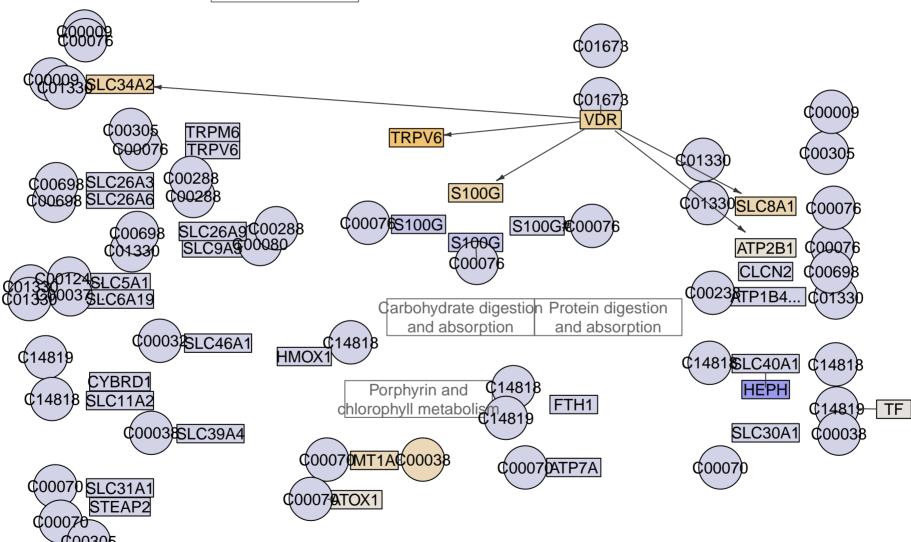




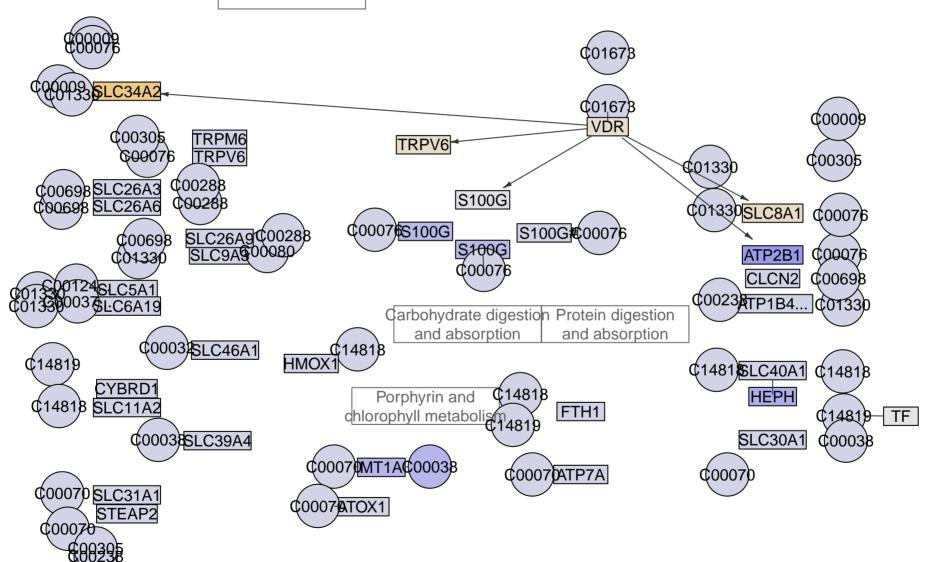


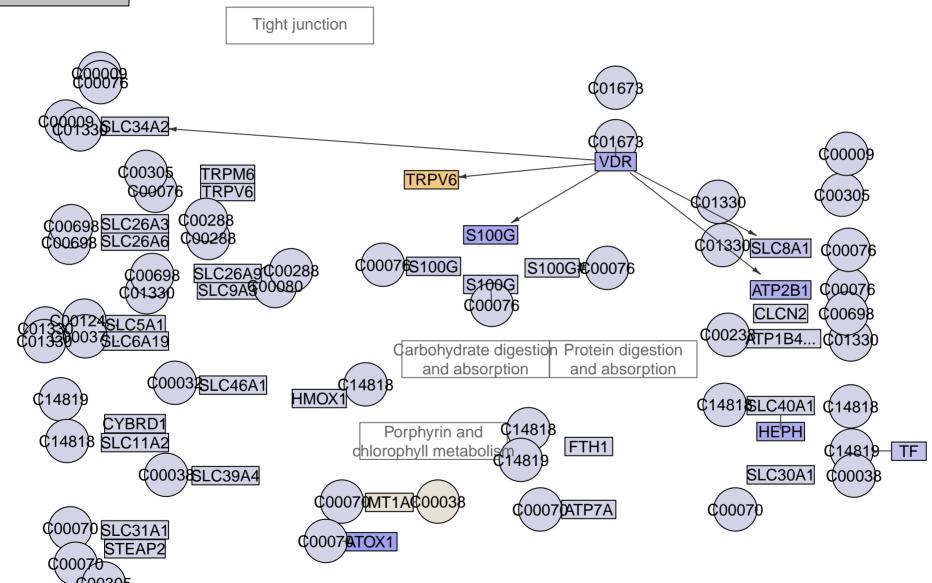


Tight junction





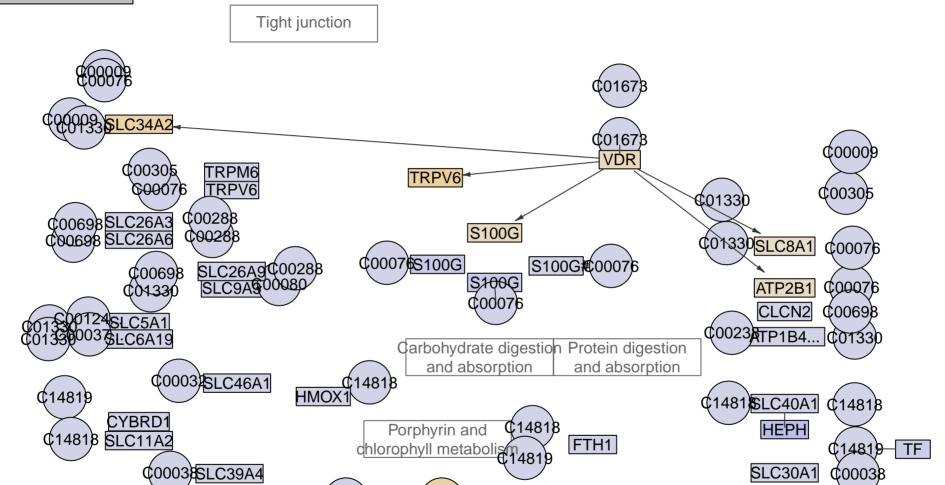




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STEAP2

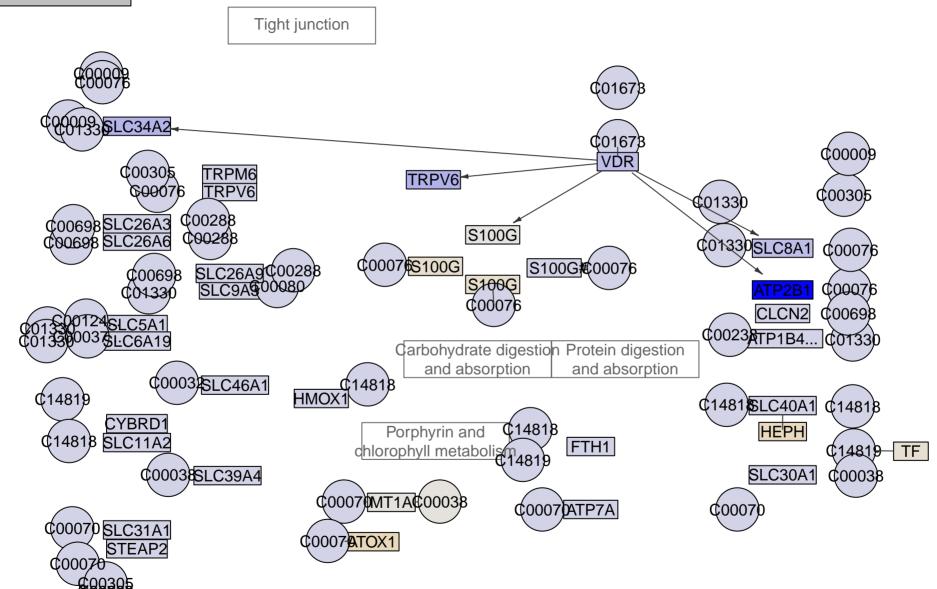


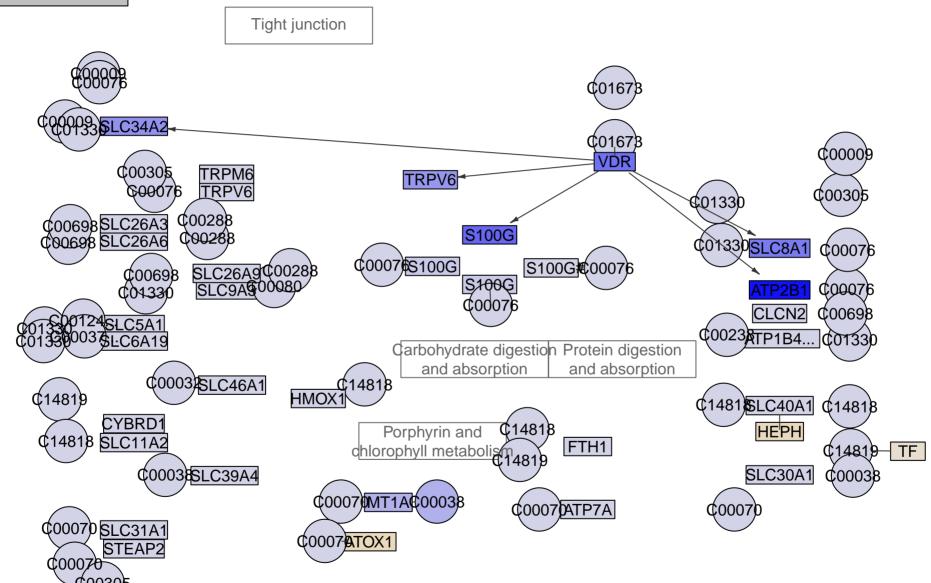
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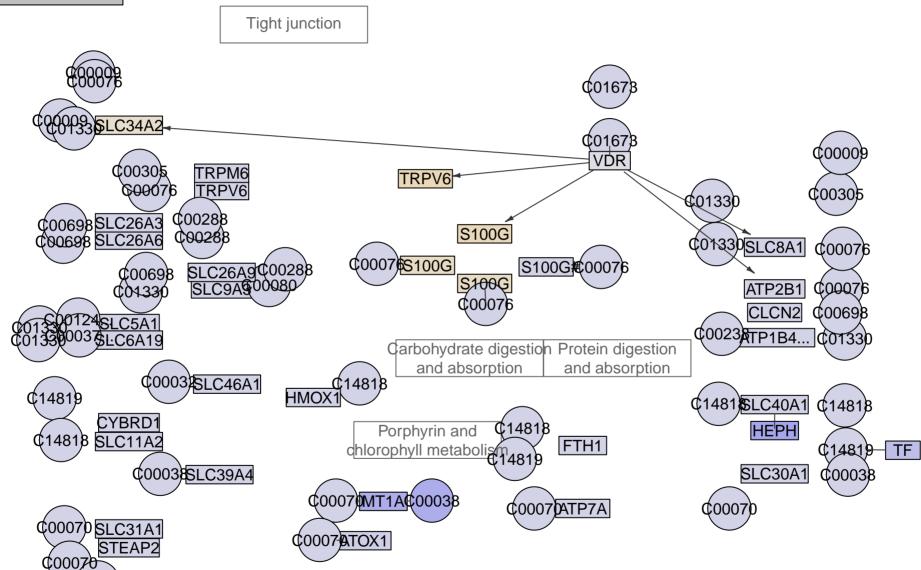
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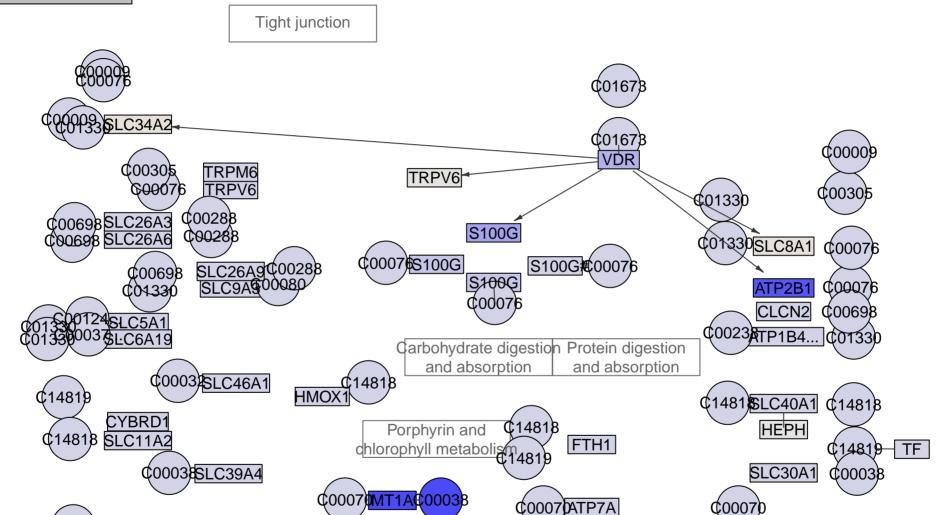




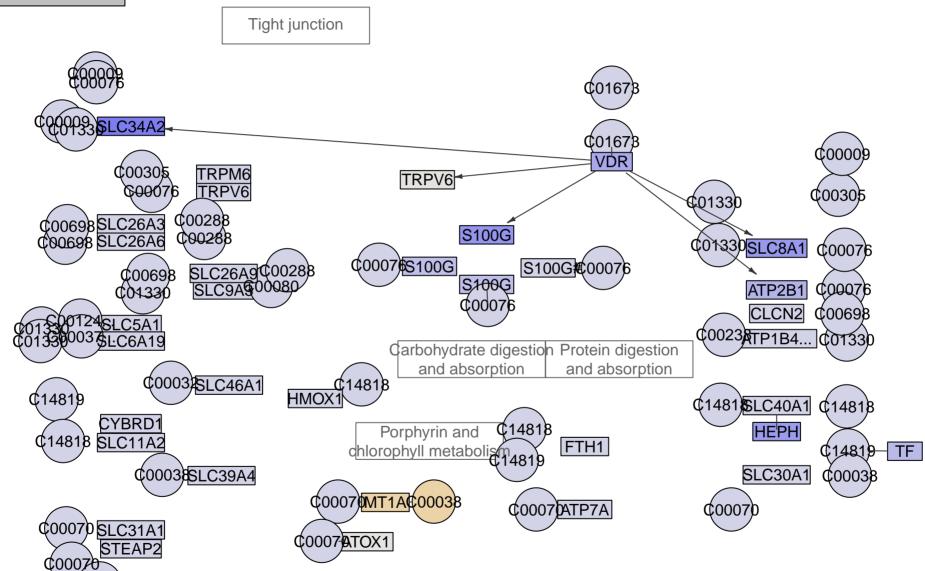
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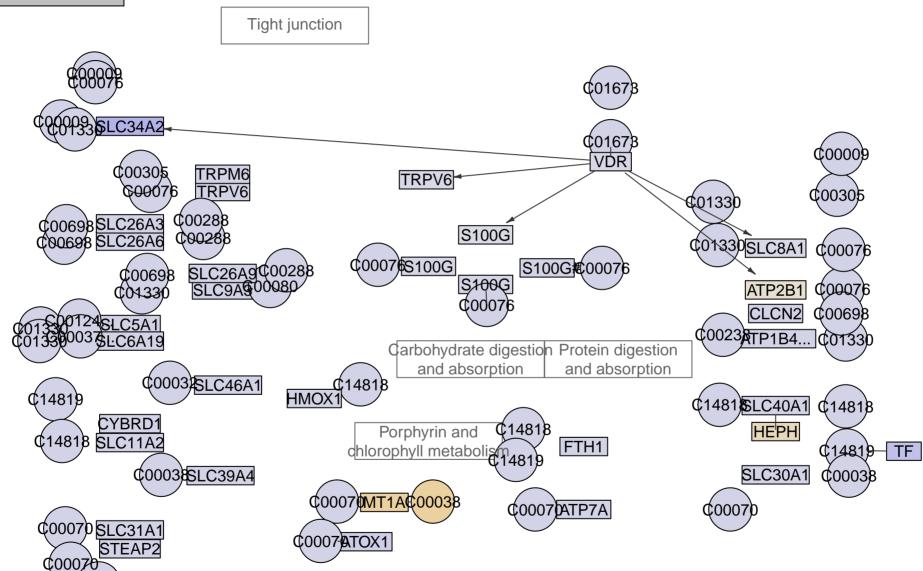
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STEAP2



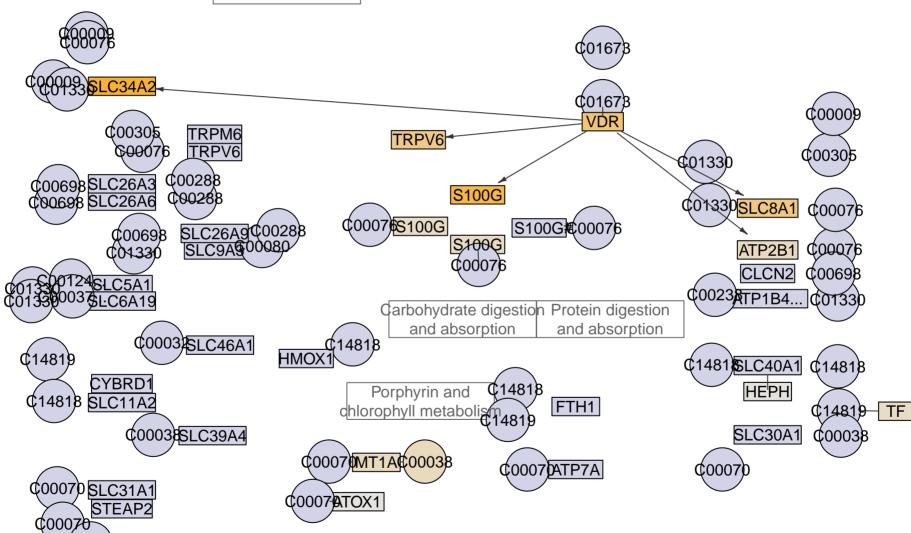
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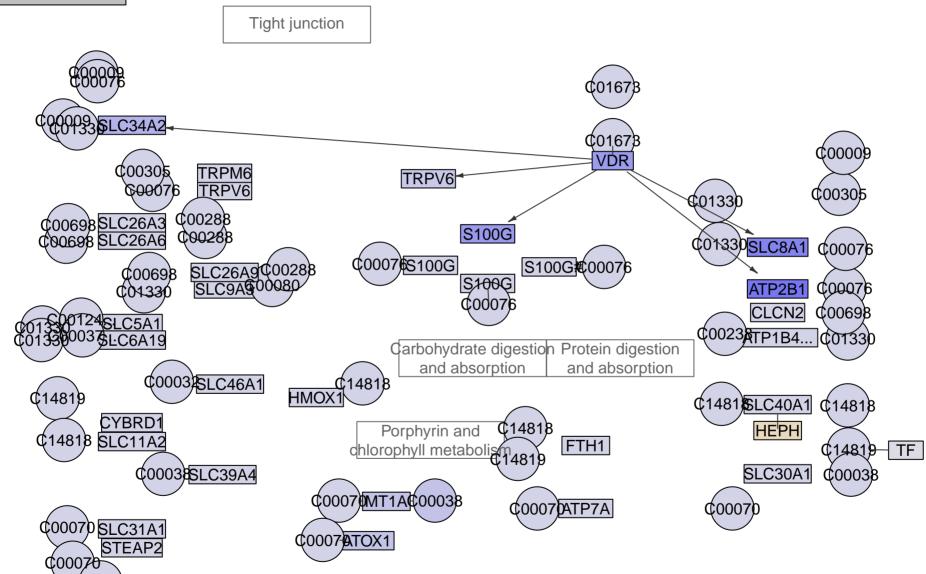


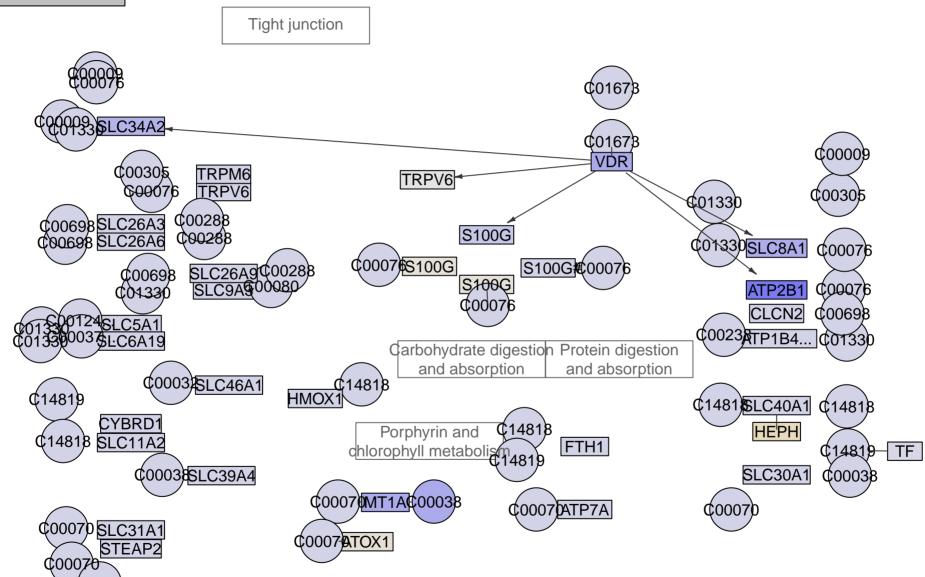


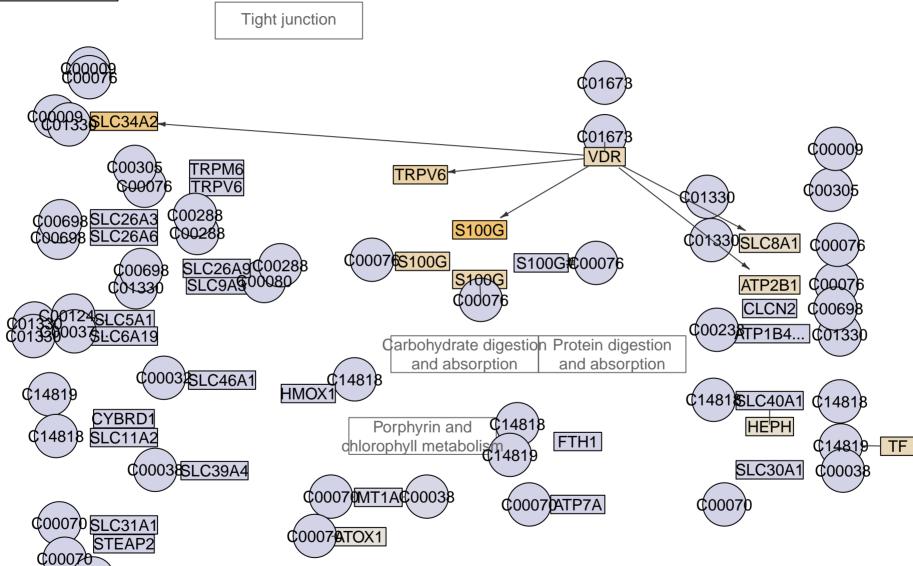
Mineral absorption
MPI–180

Tight junction

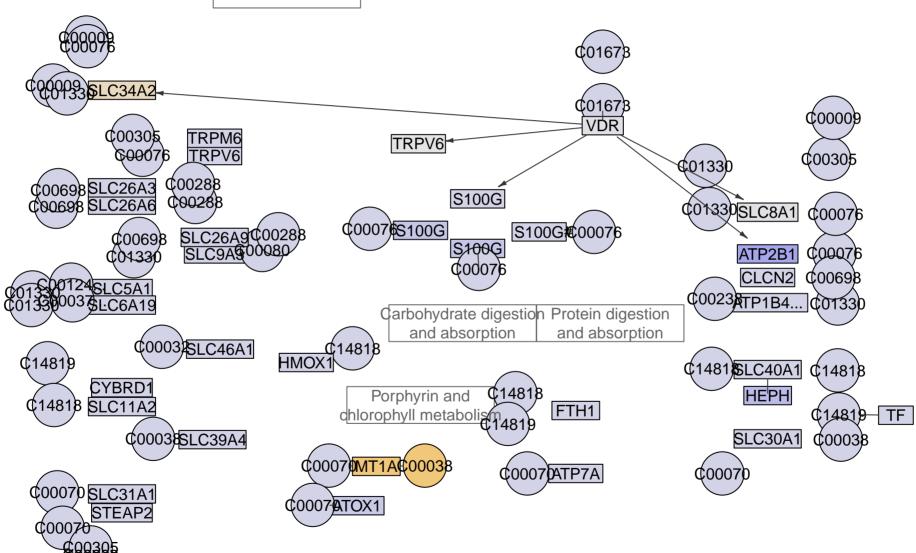


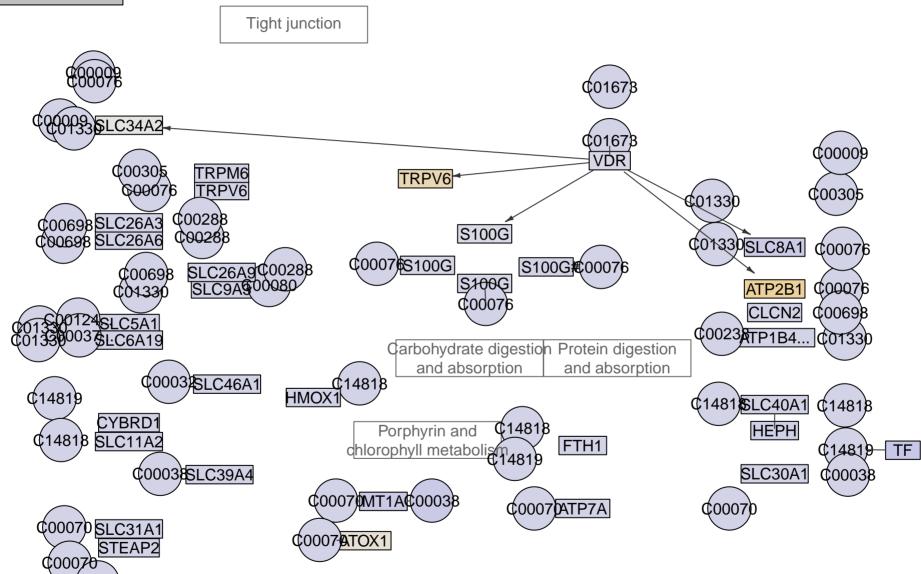


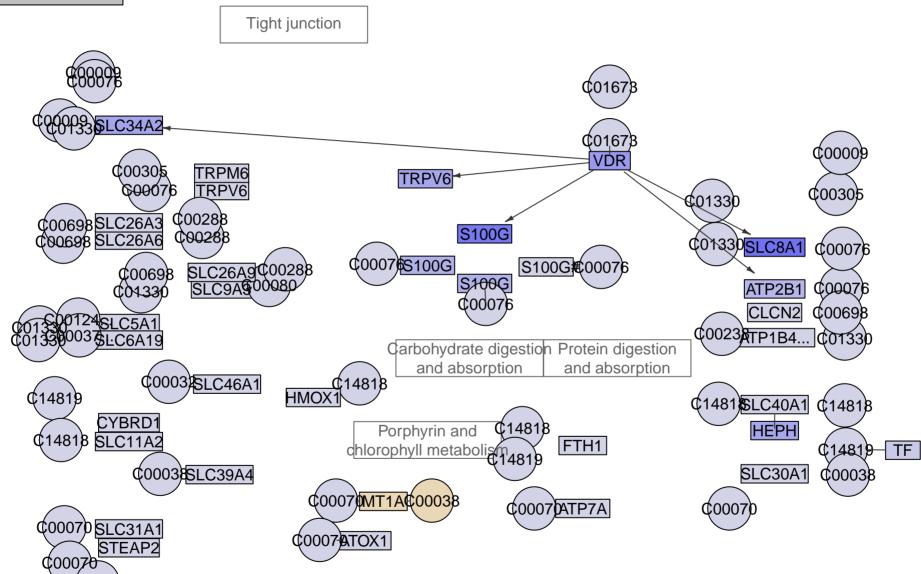


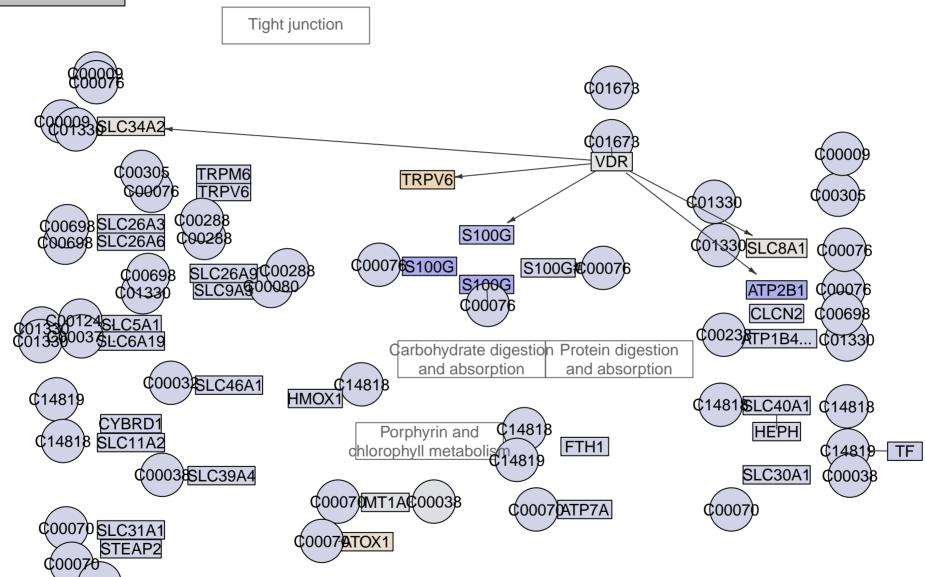


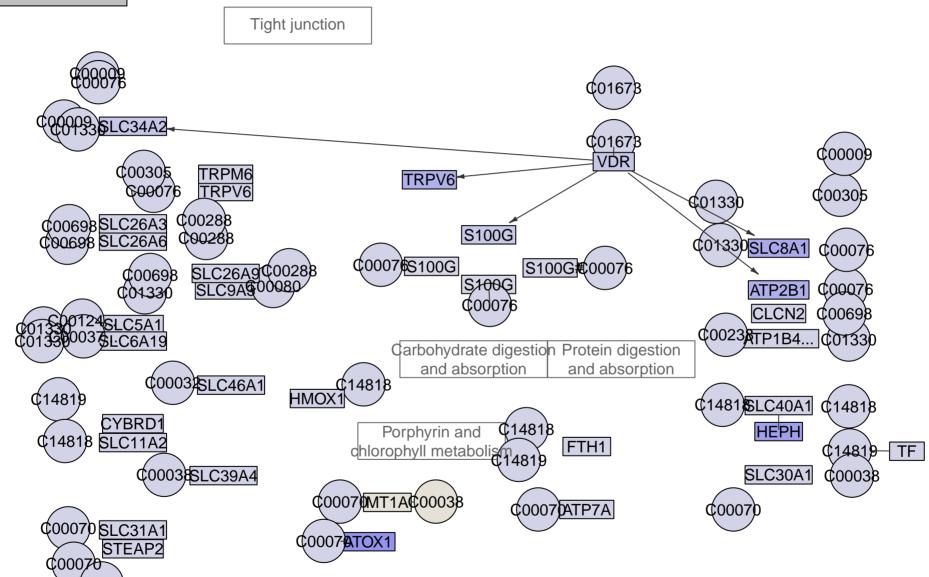






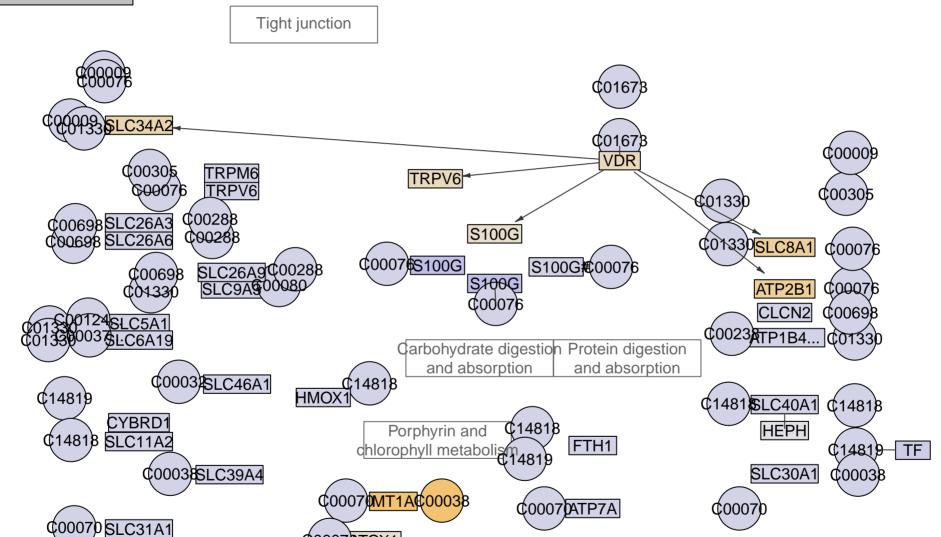




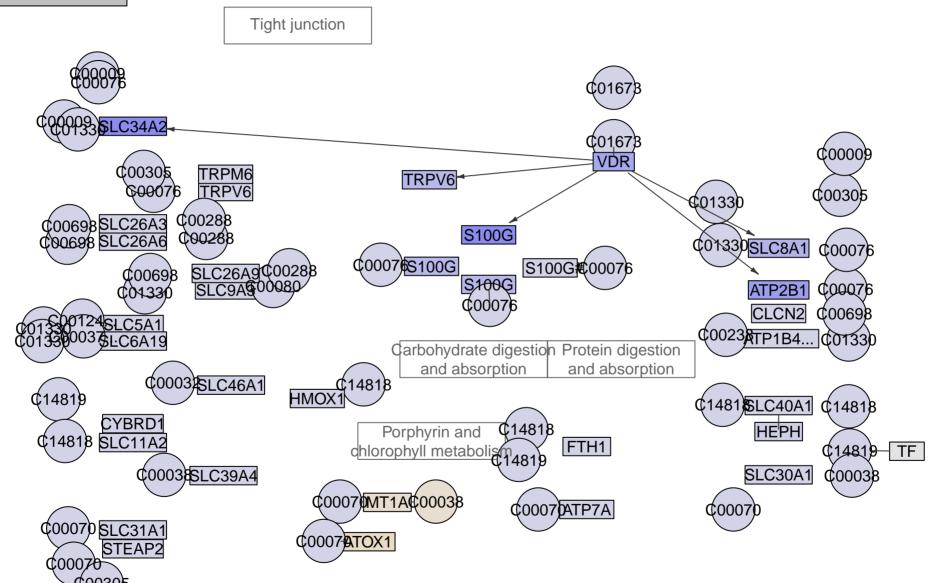


STEAP2

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**C**0007 ATOX1

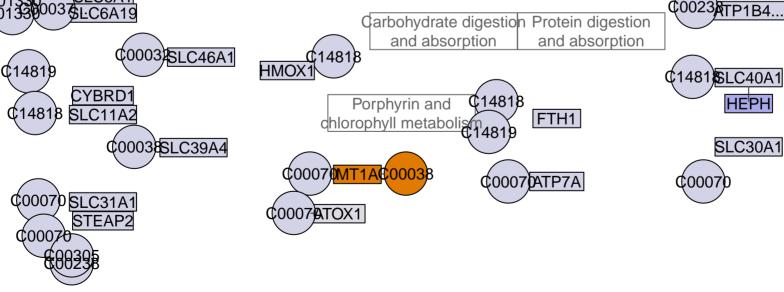


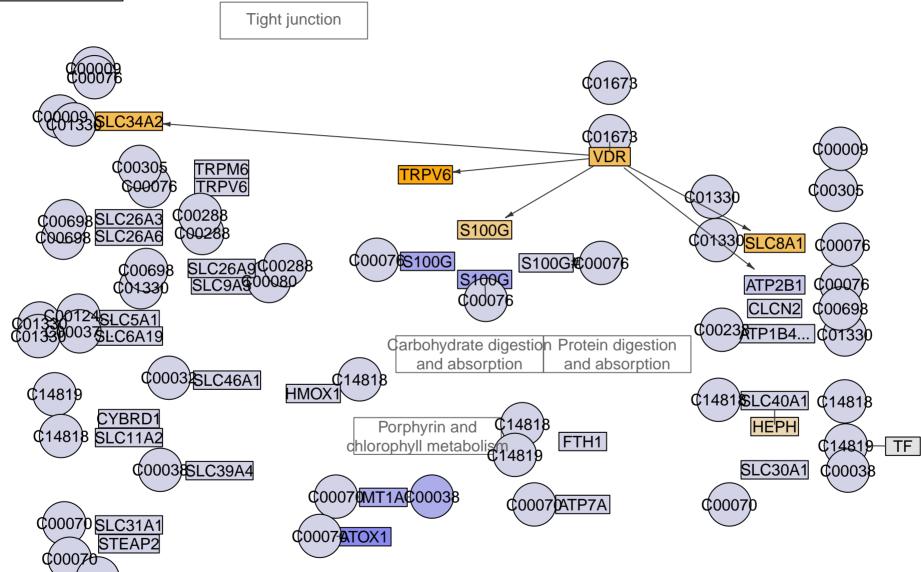
Tight junction **¢**0167 000033 SLC34A2 **C**0167 ¢0000 VDR C0030 TRPM6 TRPV6 TRPV6 600076 ¢00305 ¢01330 ¢00288 C00698 SLC26A3 C00698 SLC26A6 S100G ¢90288 (0133)SLC8A1 ¢00076 C0007 \$100G SLC26A9(0028) SLC9A(00080 S100G#00076 ¢00698 S100G ATP2B1 C00076 **¢**04330 C00076 CLCN2 C00698 00124SLC5A1 ¢0023 003 \$LC6A19 C01330 Carbohydrate digestion Protein digestion and absorption and absorption C0003 SLC46A1 HMOX C1481 SLC40A1 C14818 CYBRD<sup>2</sup>

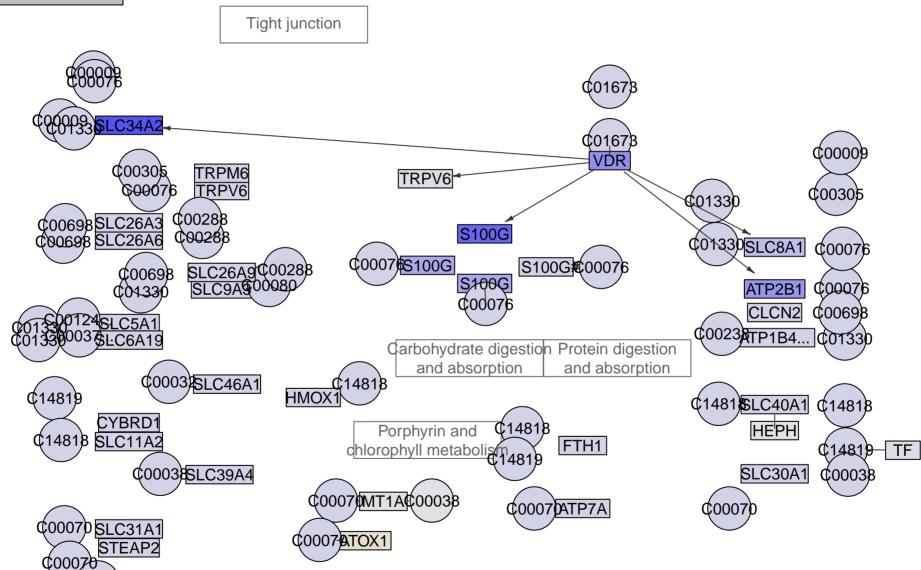
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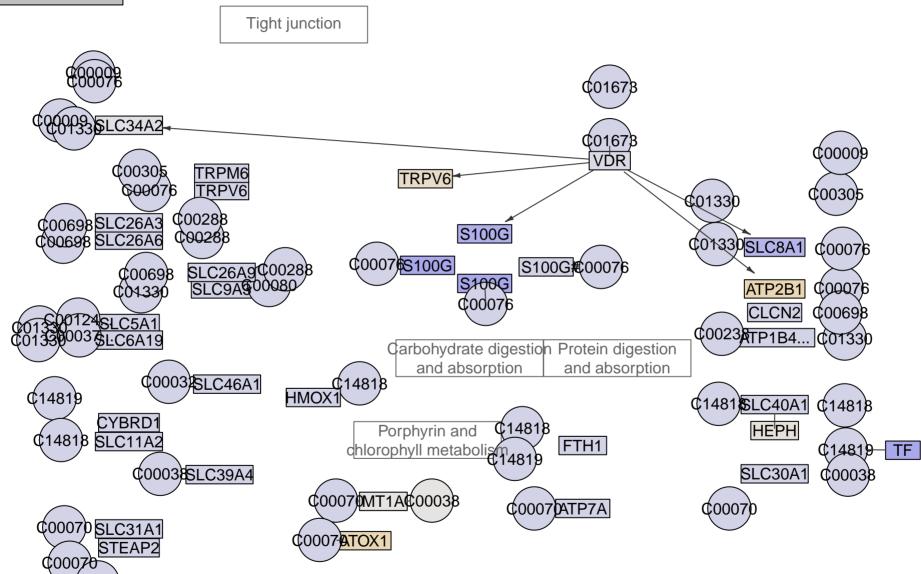
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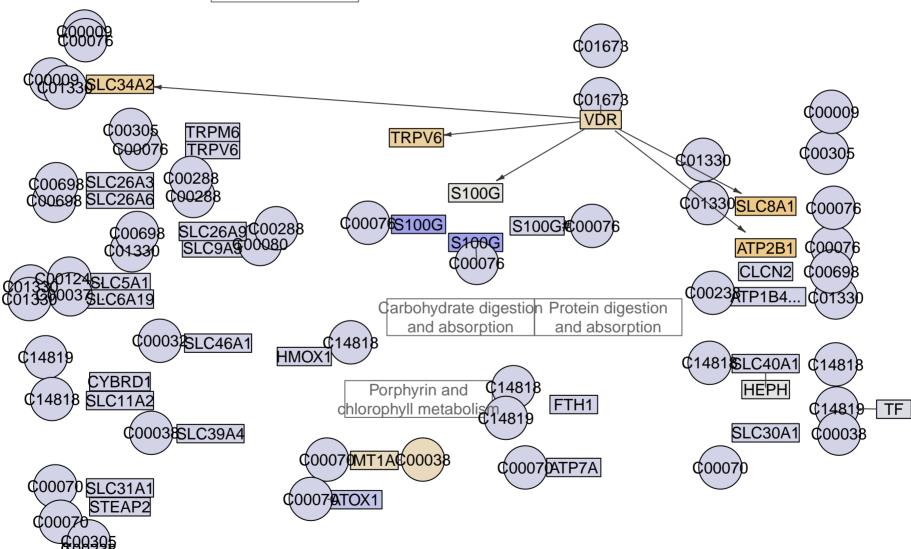








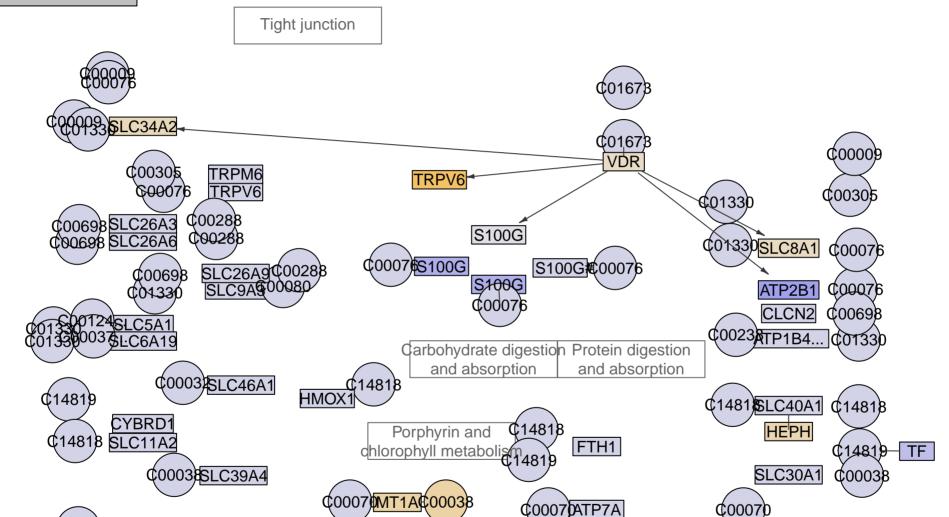




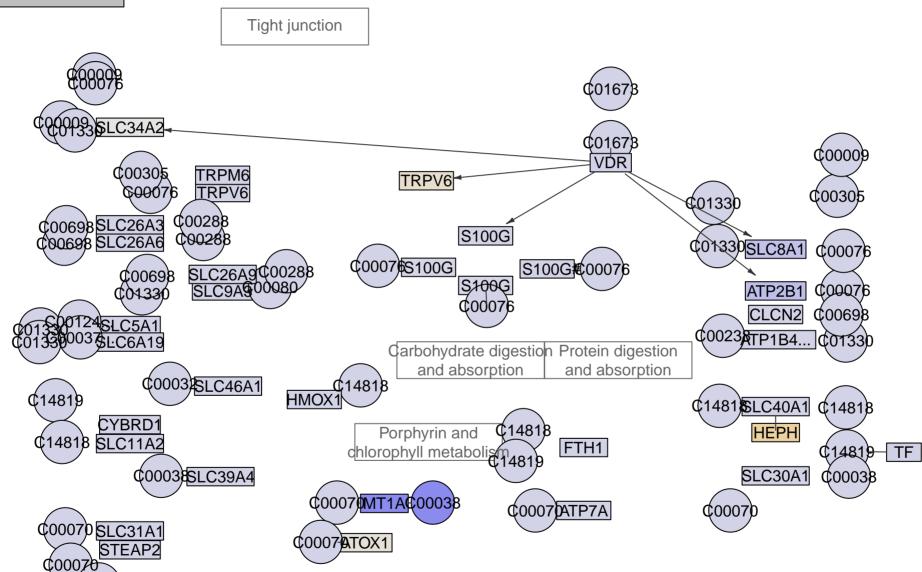
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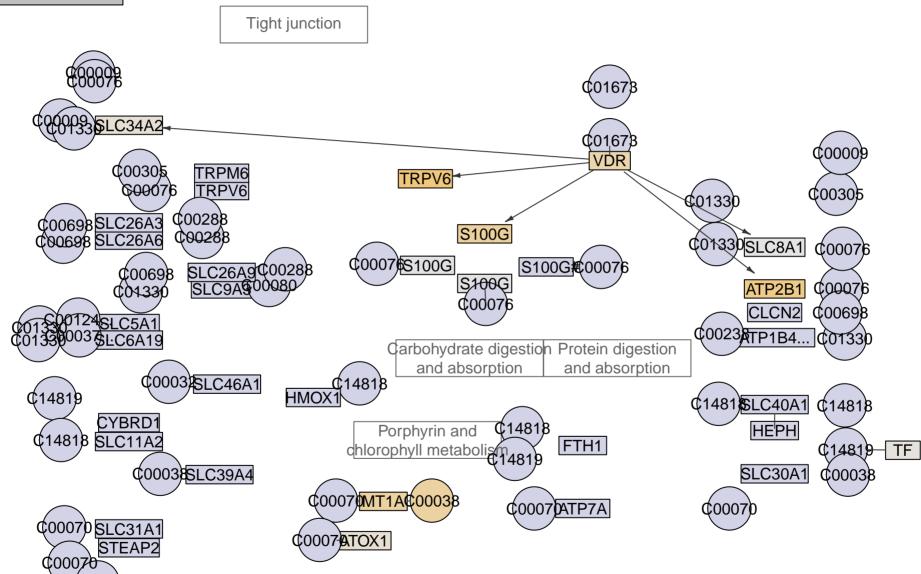
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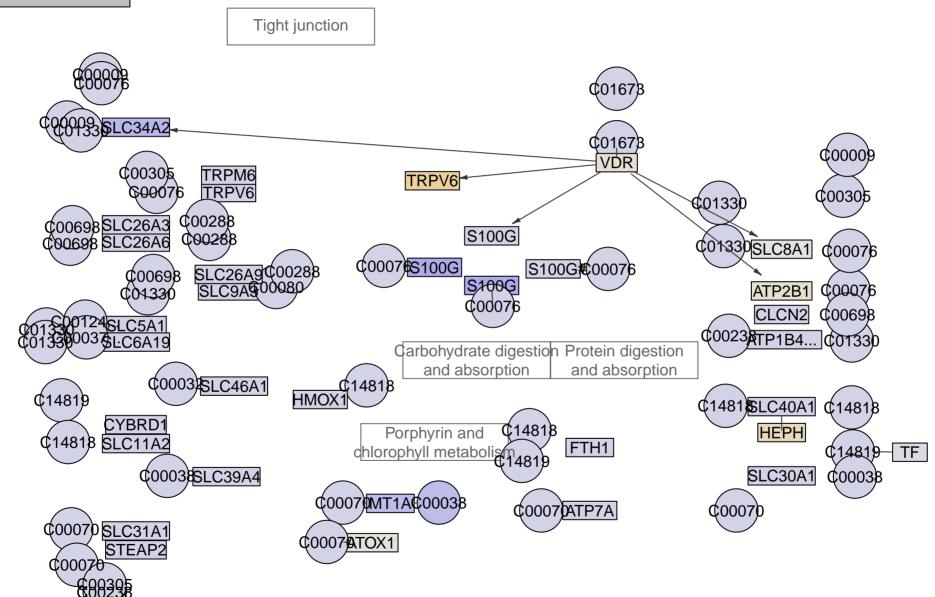
STEAP2



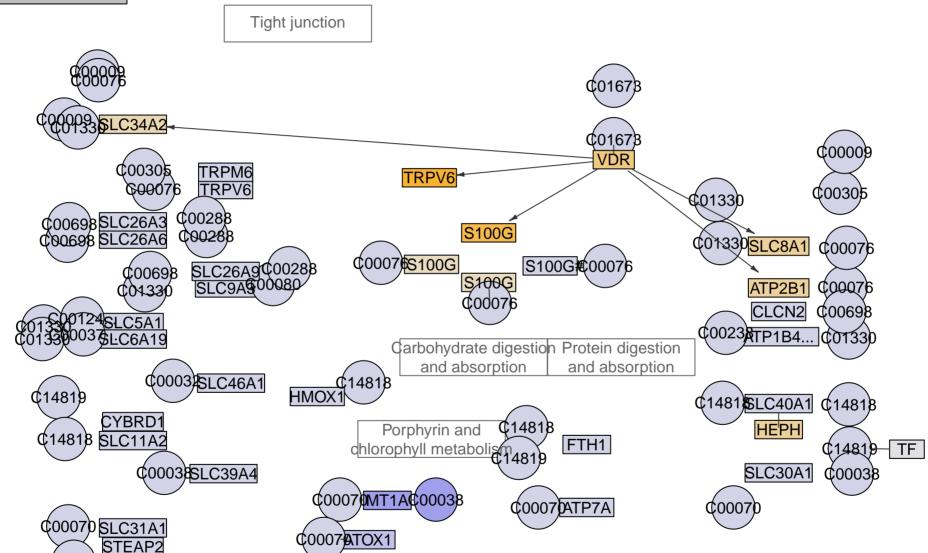
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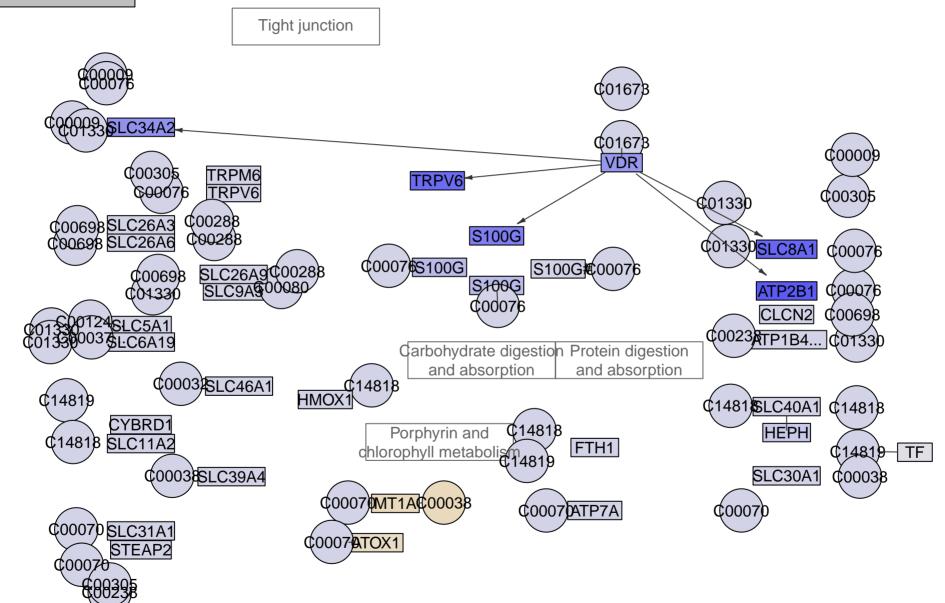


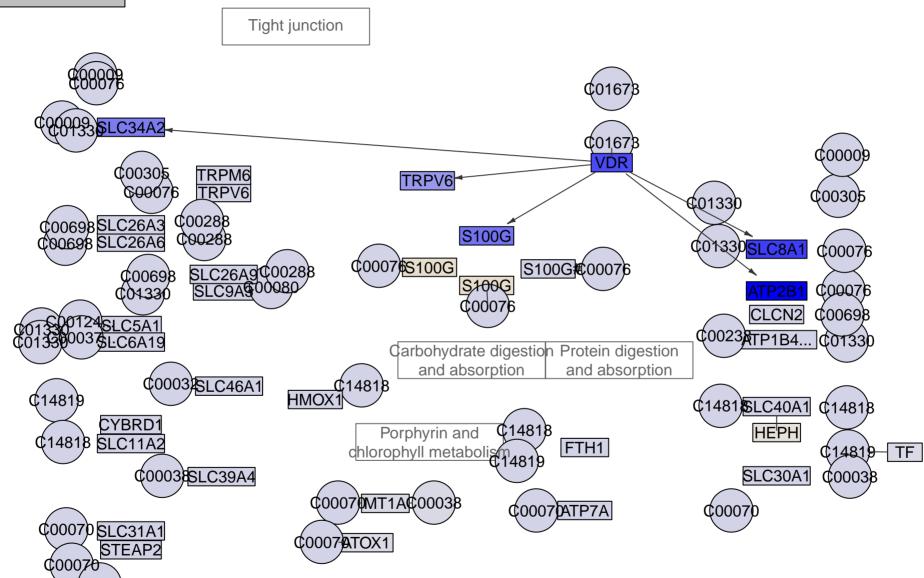


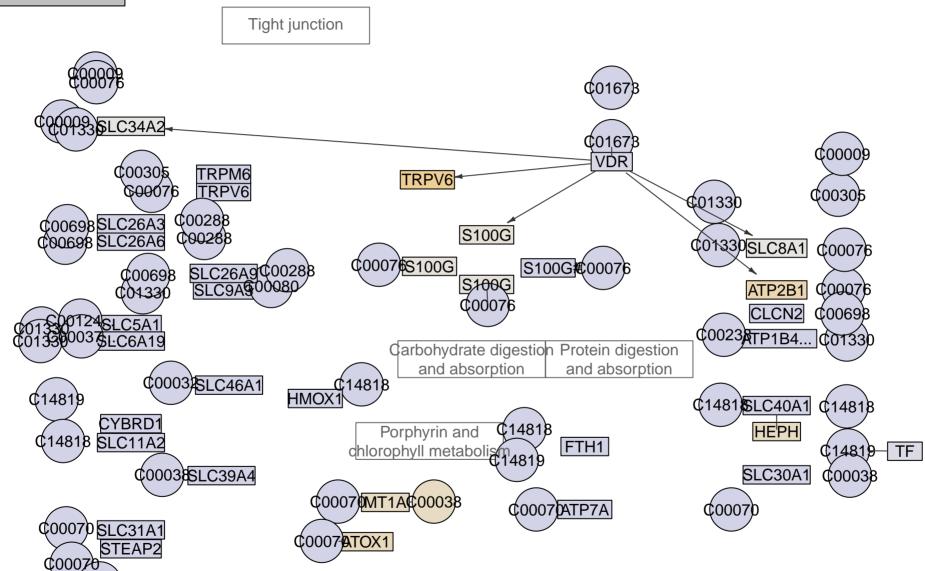


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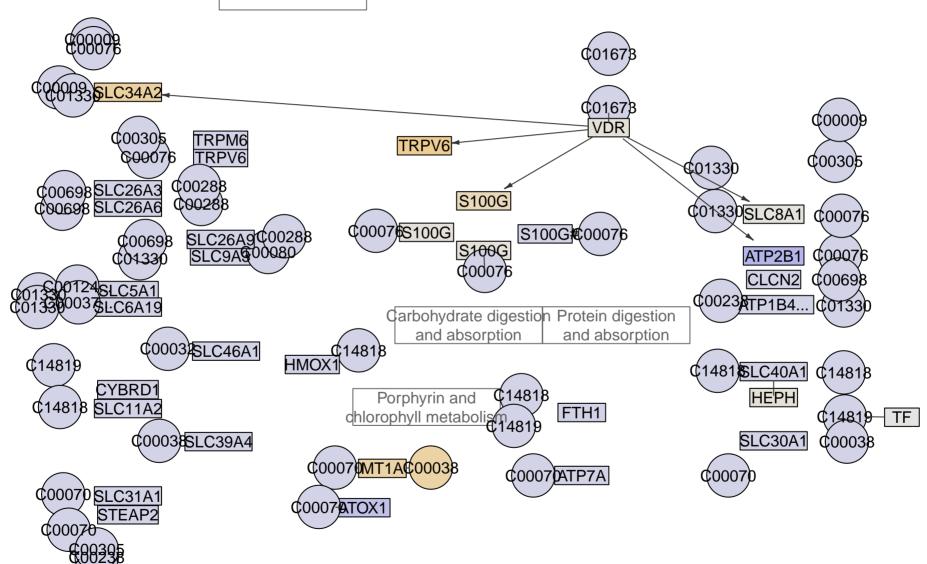


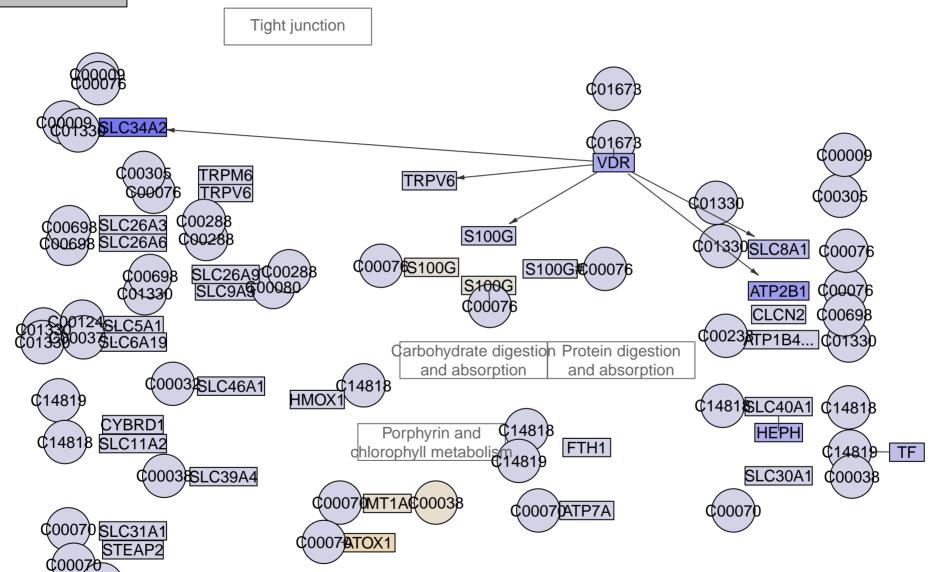


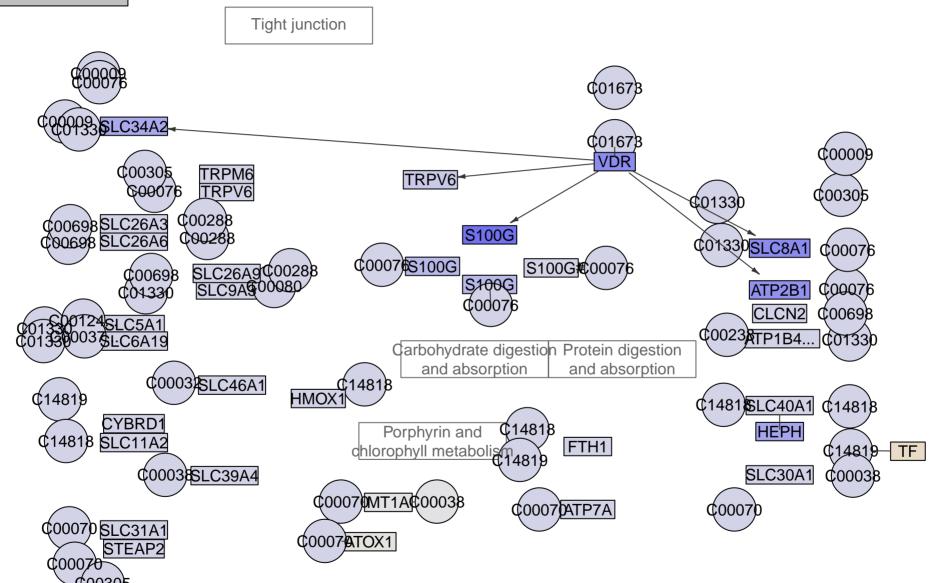


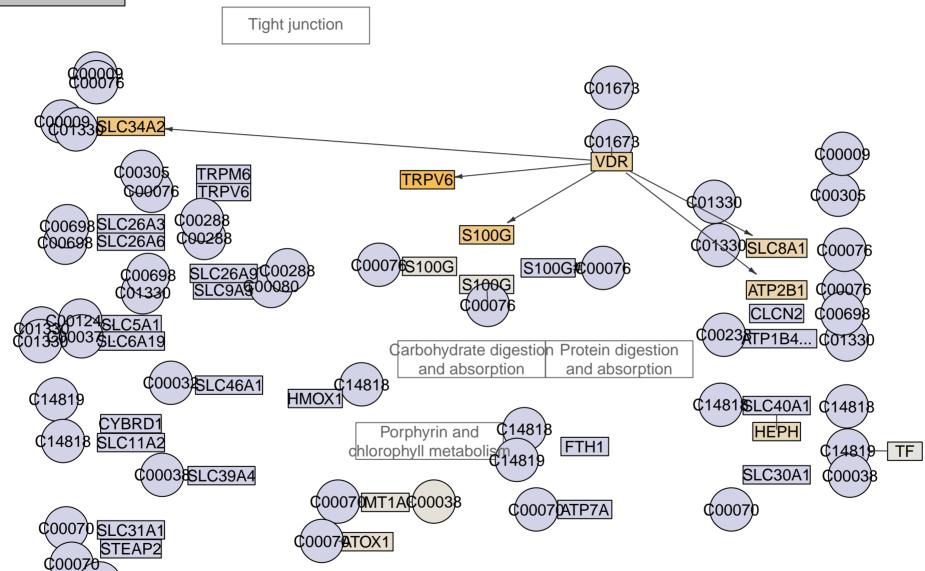


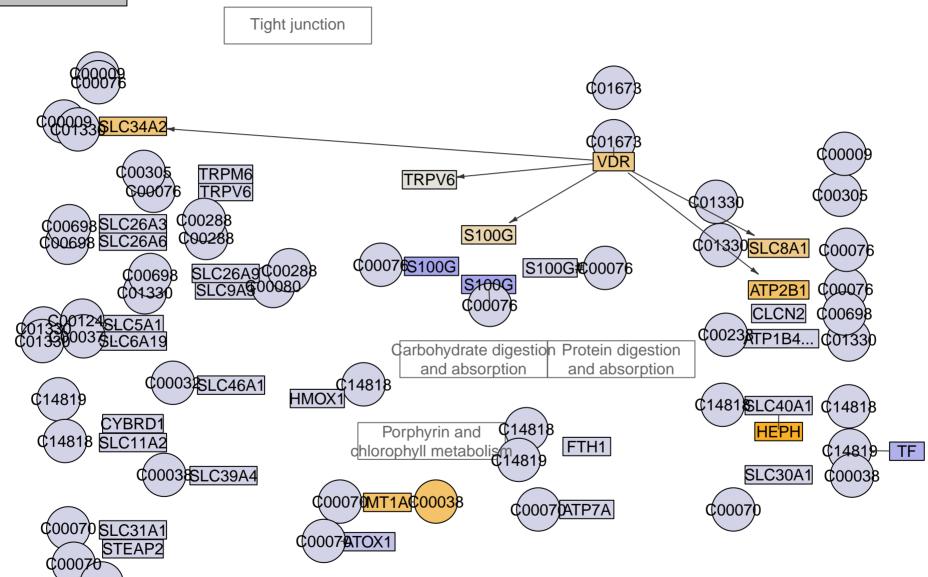
Tight junction

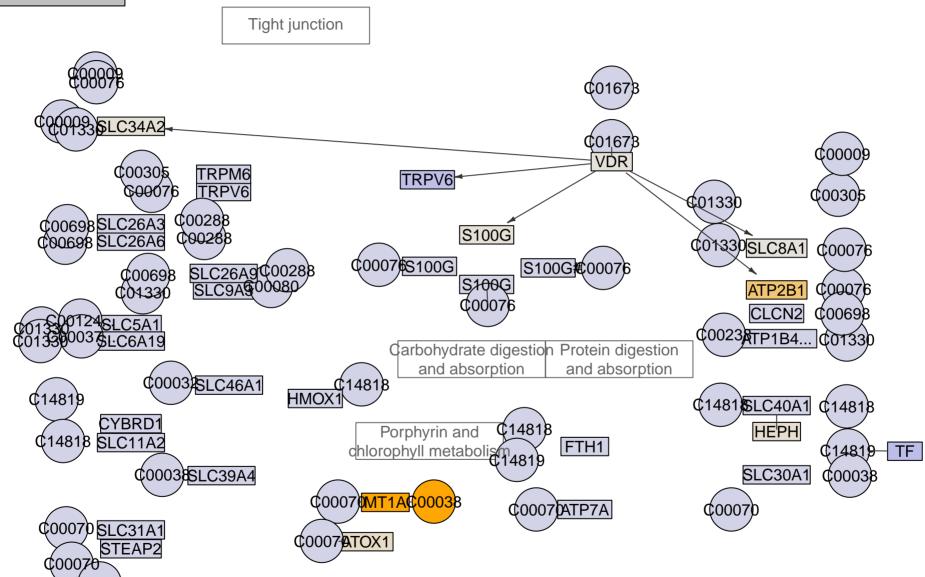


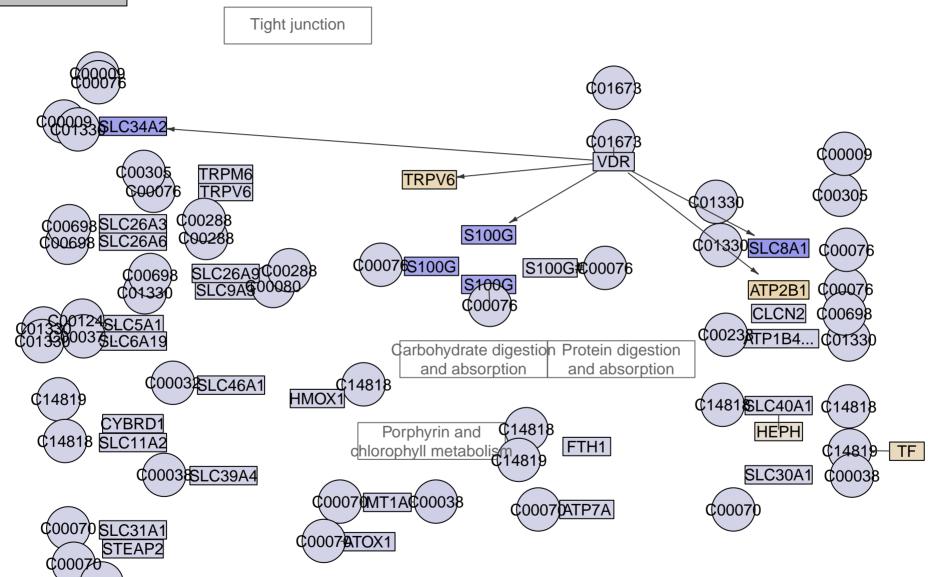


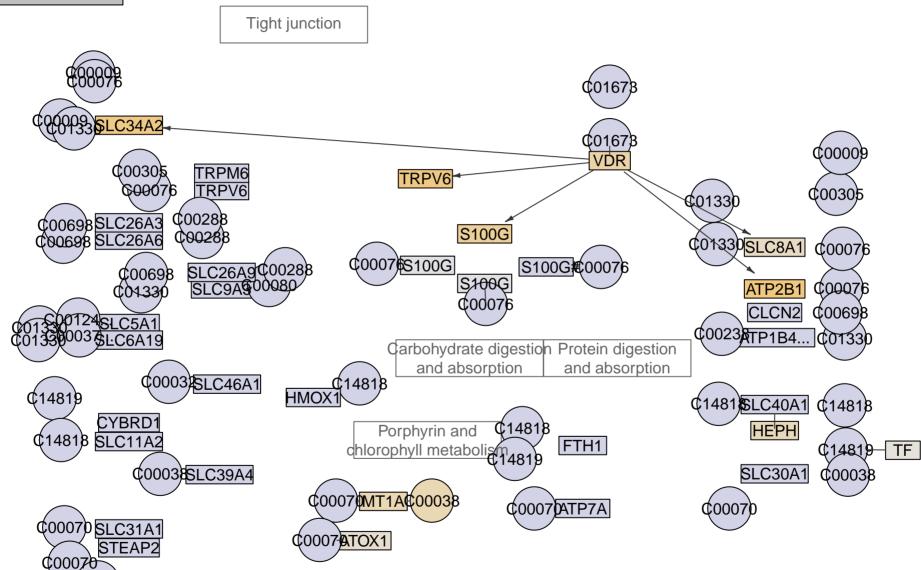


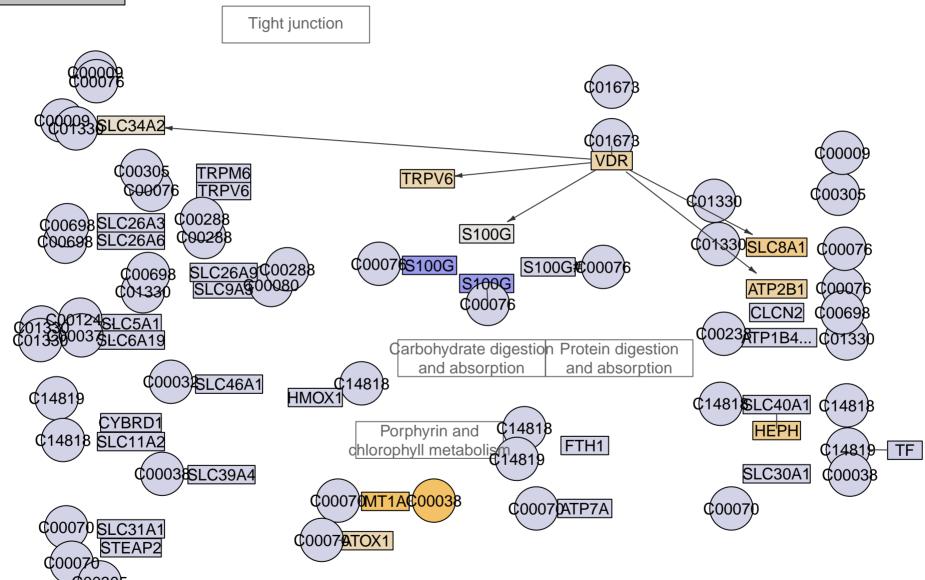


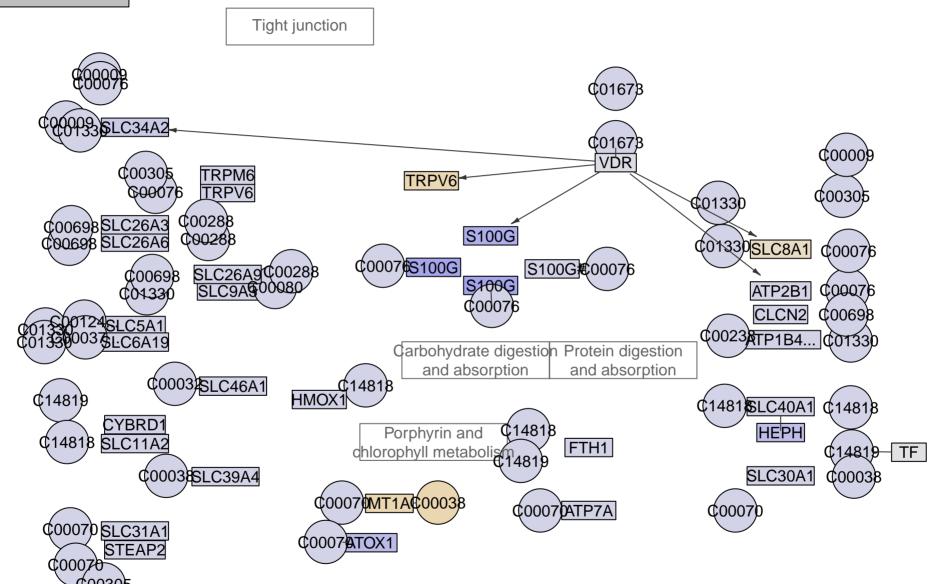




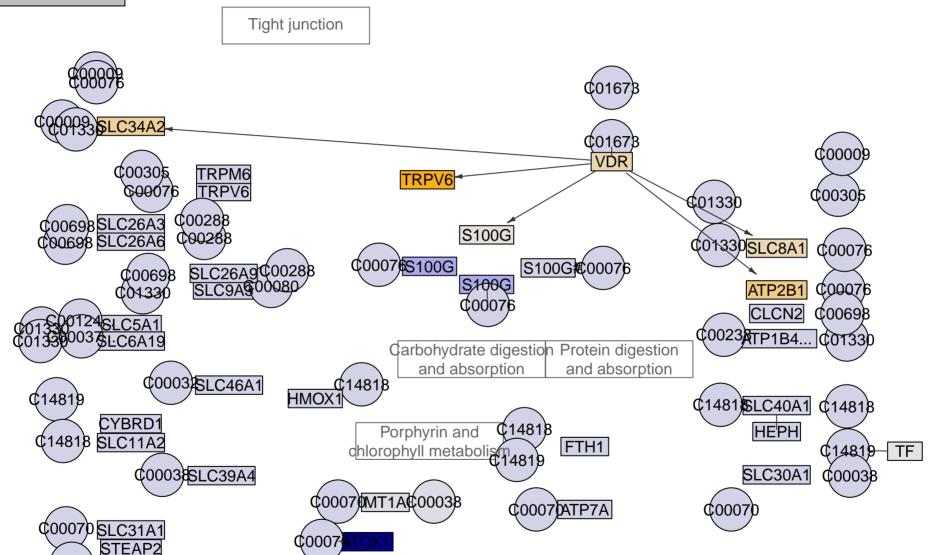


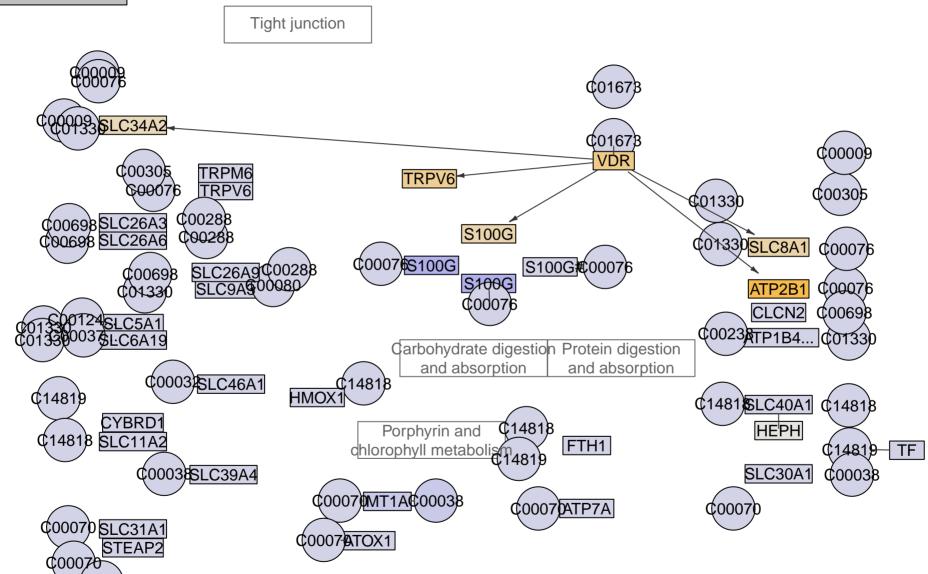


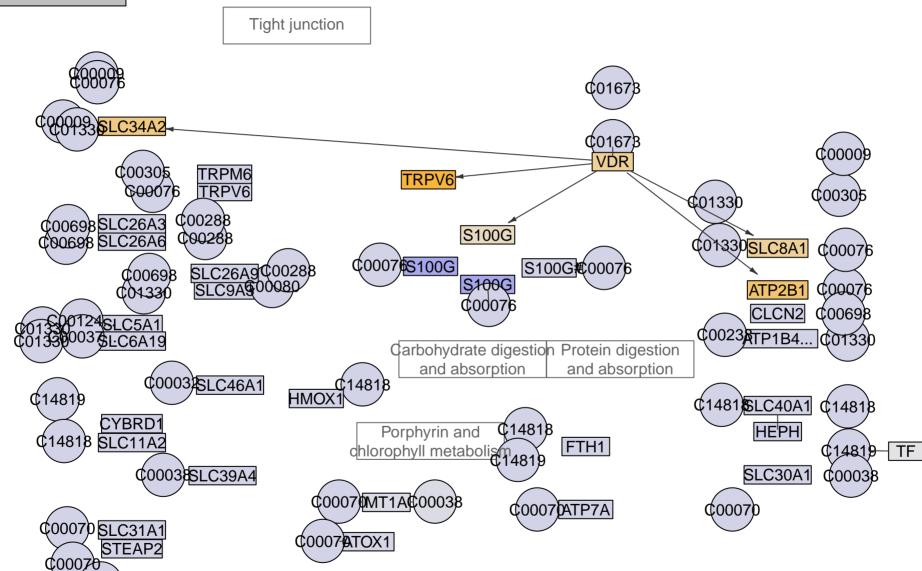




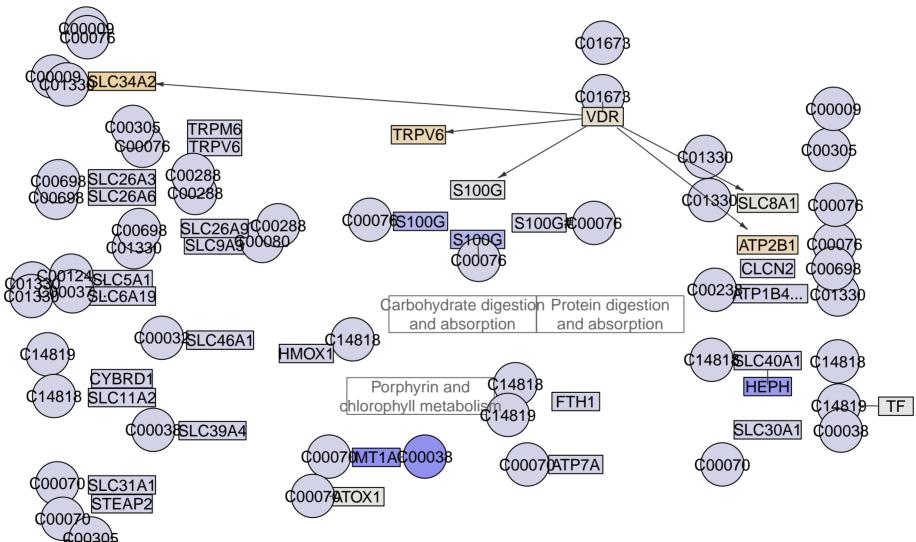
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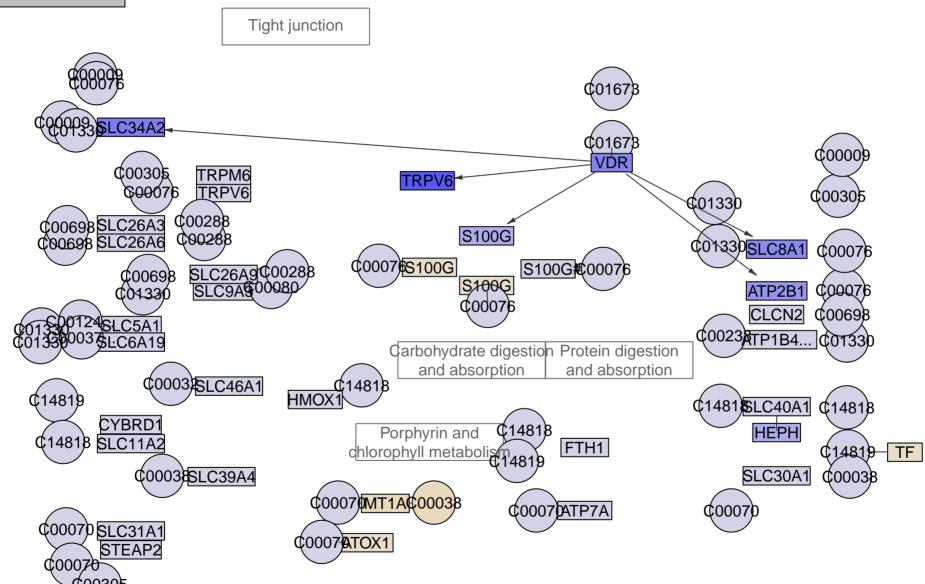


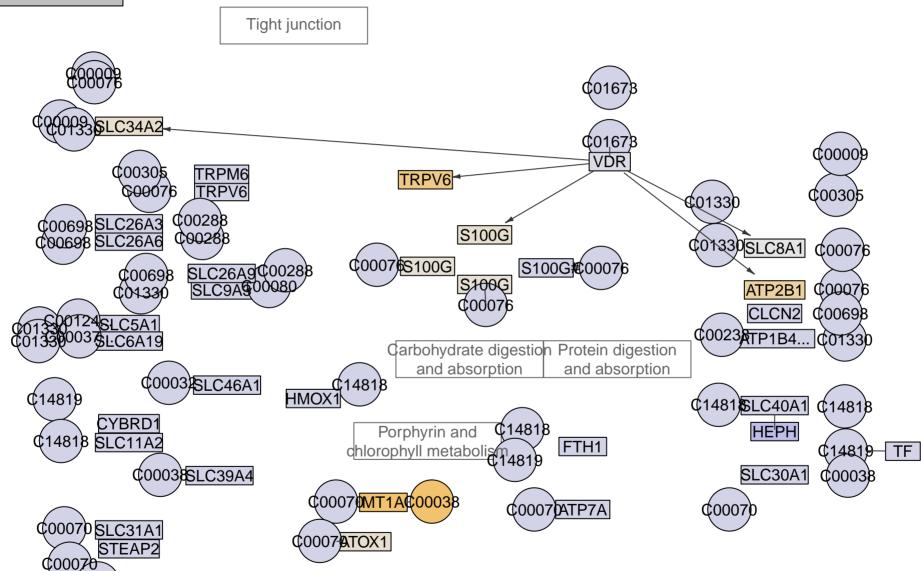


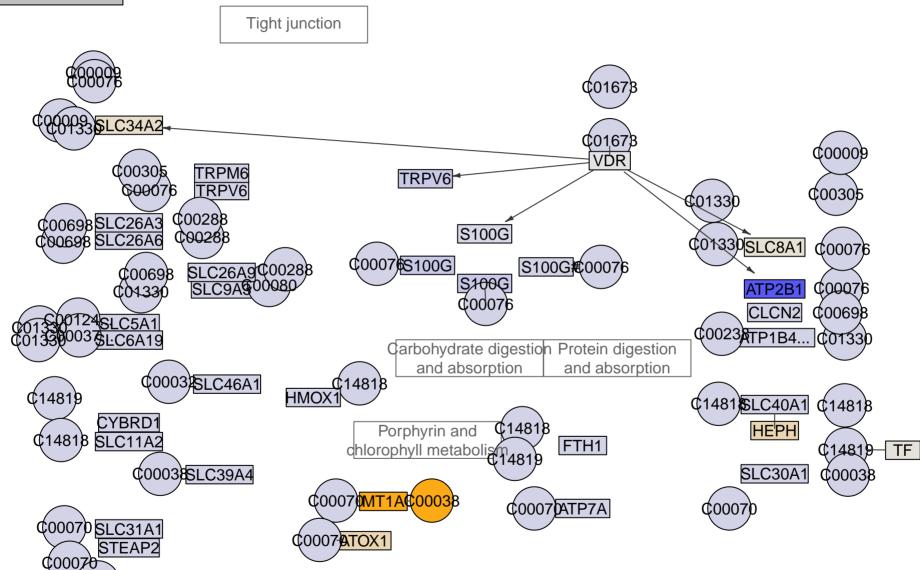


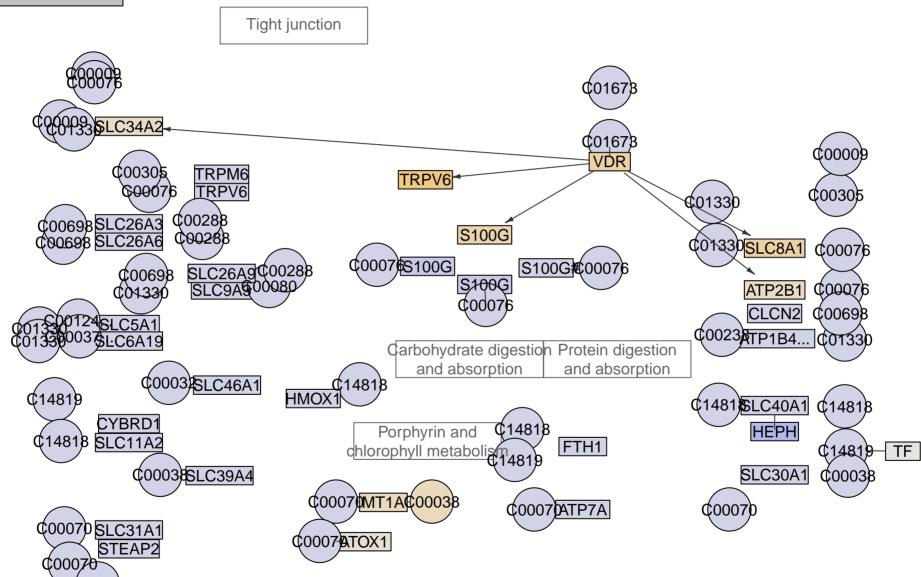
Tight junction

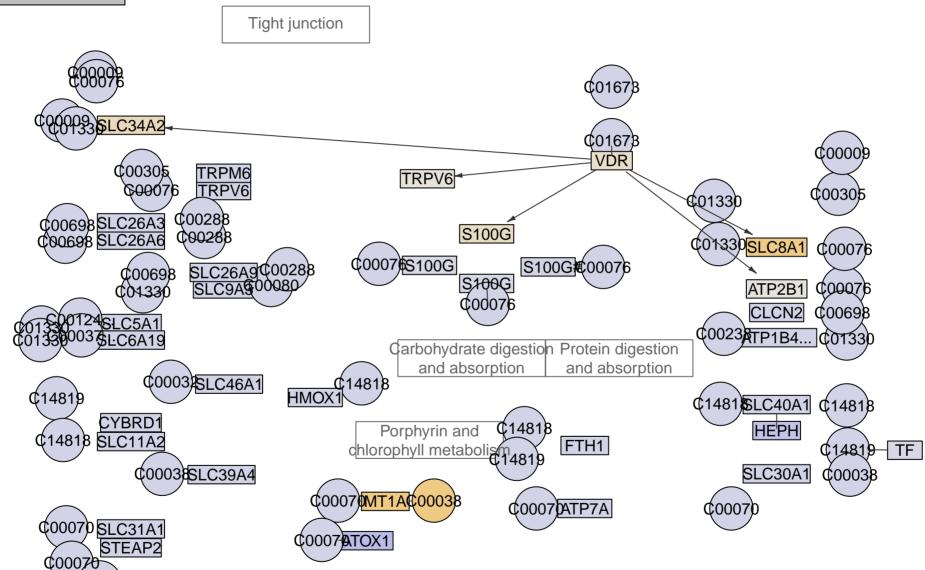


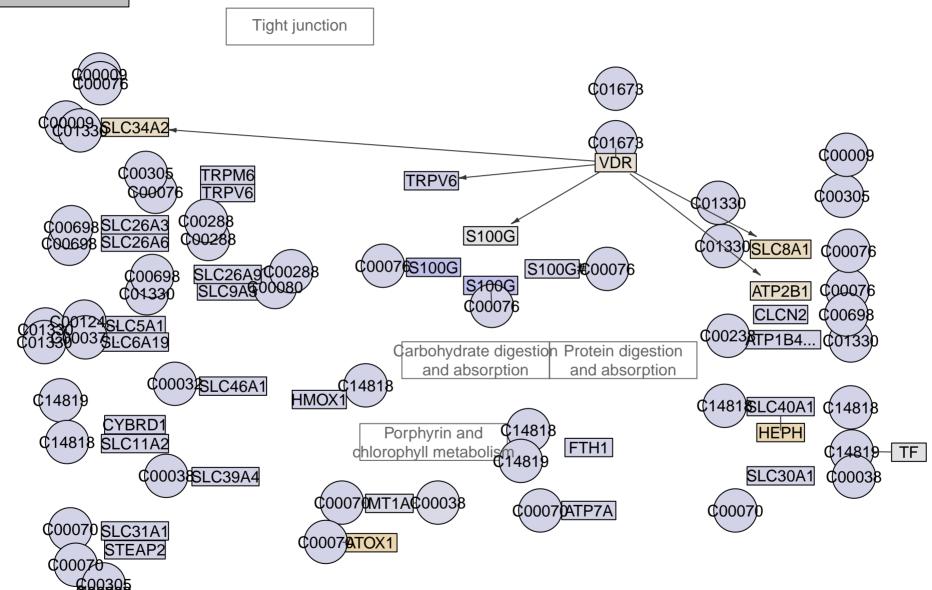


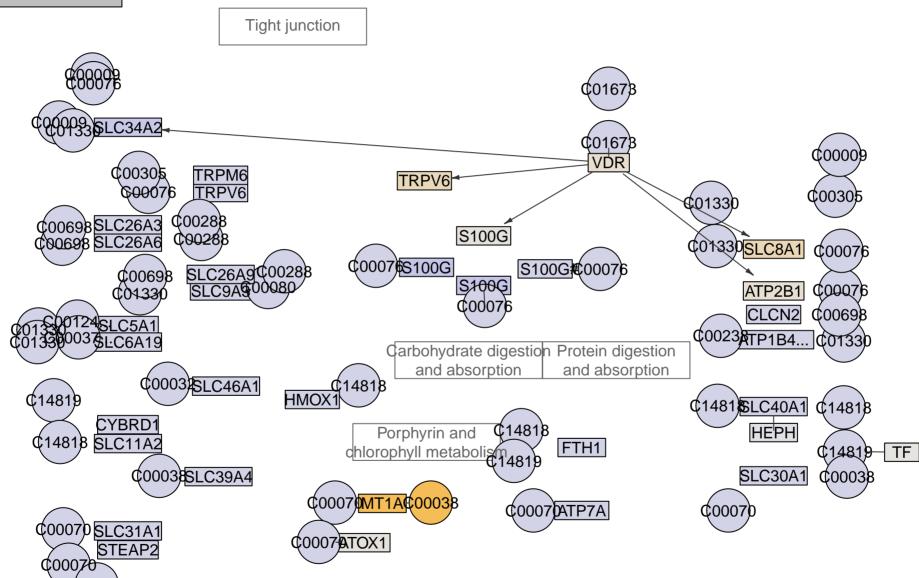


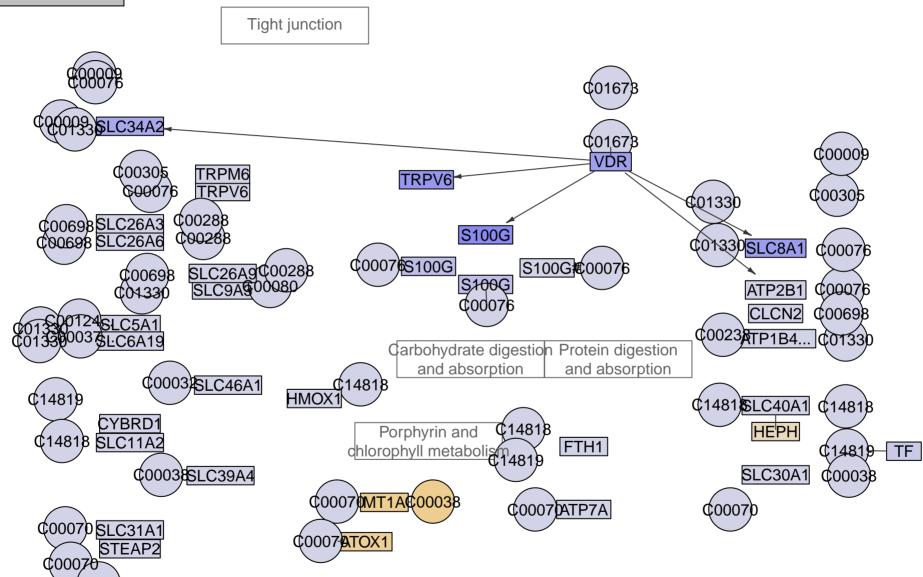


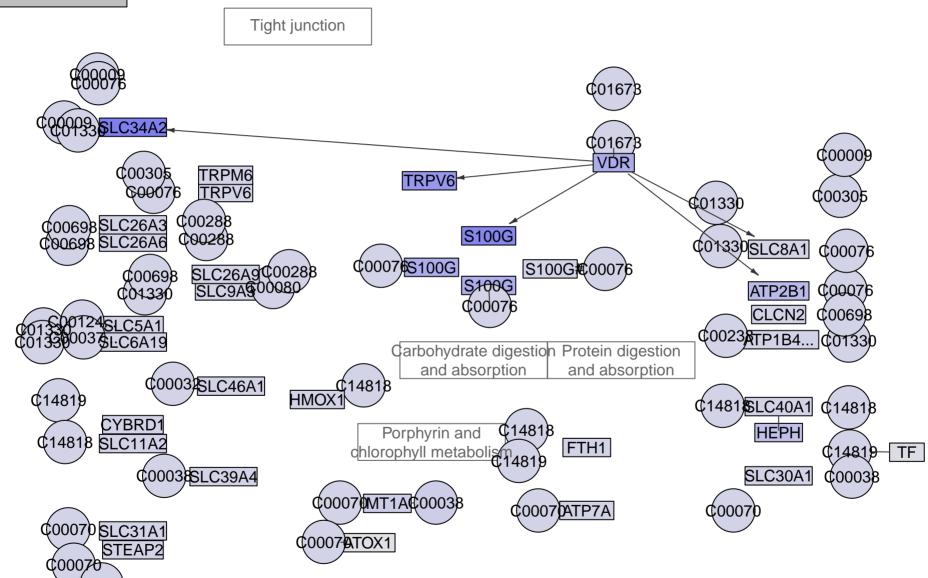






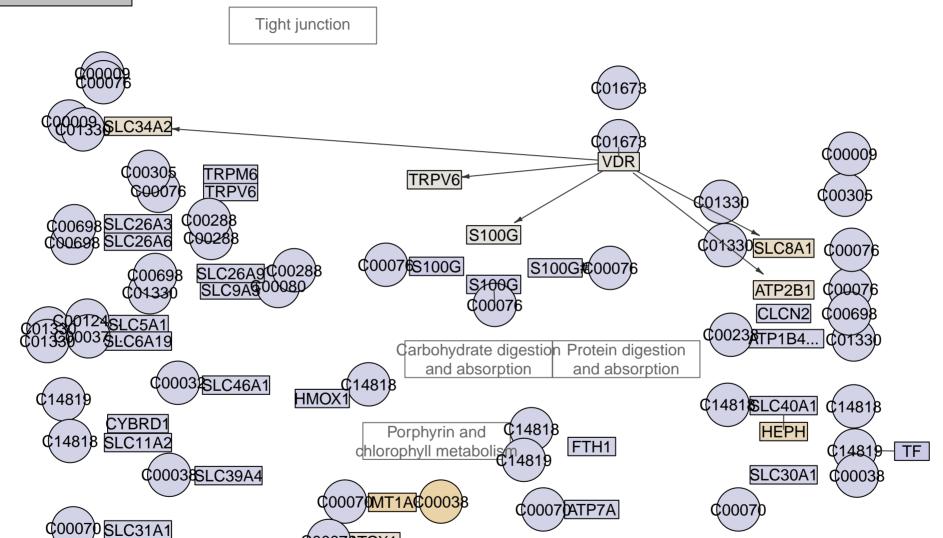






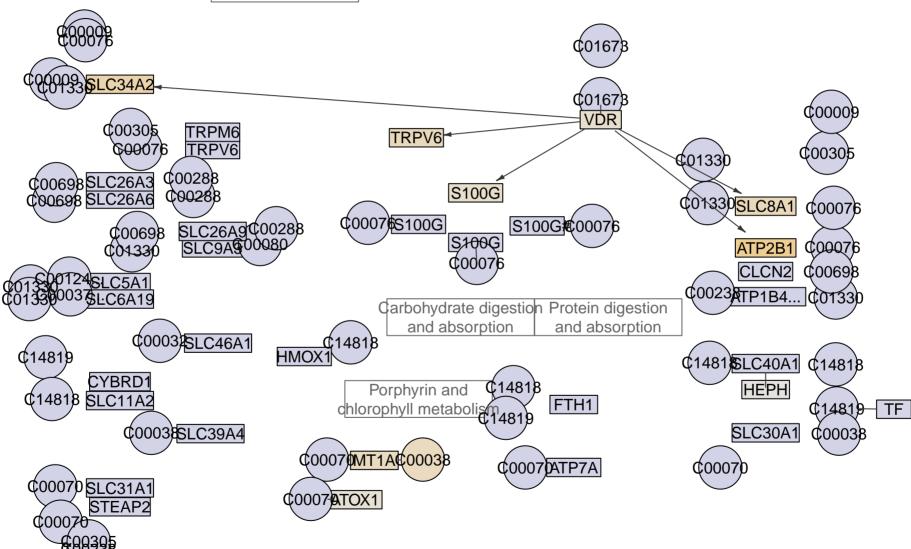
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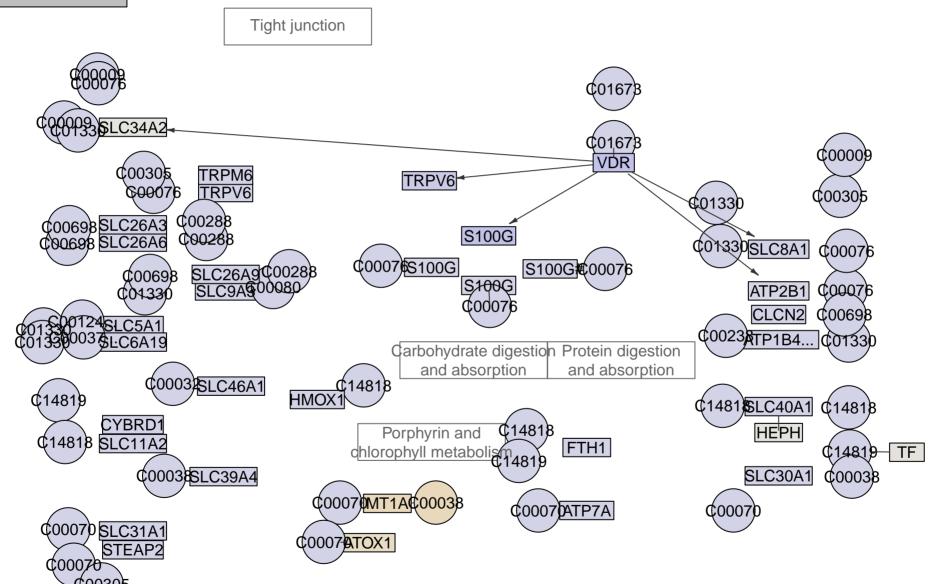
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C1481 SLC40A1 C14818 CYBRD<sup>2</sup> C14818 Porphyrin and HE<sub>PH</sub> chlorophyll metabolis (1481) C14818 SLC11A2 FTH1 ¢14819 TF C0003 SLC39A4 SLC30A1 ¢00038 C00070MT1AC00038 C0007)ATP7A (0007) ¢00070 SLC31A1

**C**0007 ATOX1



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