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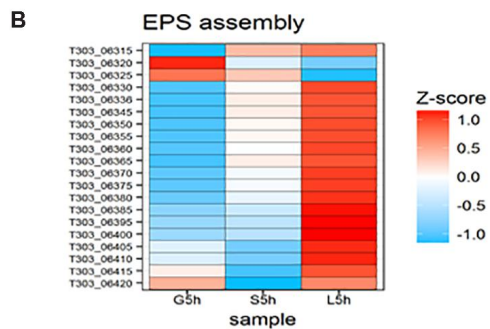
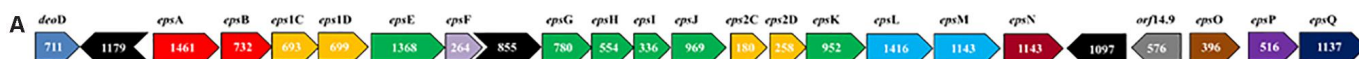
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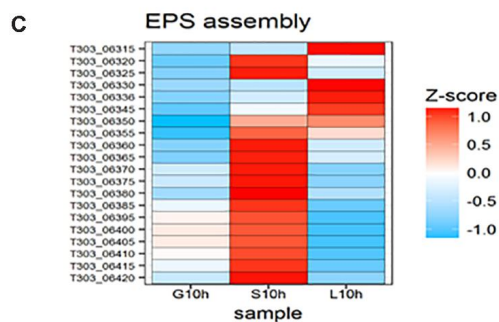
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Gene_id	Gene_annotation	Function
T303_06315	<i>epsQ</i>	Permease
T303_06320	<i>epsP</i>	Phosphoglycerate mutase
T303_06325	<i>epsO</i>	Phosphatase
T303_06330	<i>orf14.9</i>	Regulation
T303_06336	<i>epsN</i>	UDP galactopyranose mutase
T303_06345	<i>epsM</i>	Polymerization & translocation
T303_06350	<i>epsL</i>	Polymerization & translocation
T303_06355	<i>epsK</i>	Glycosyl transferase
T303_06360	<i>eps2D</i>	Chain length determination
T303_06365	<i>eps2C</i>	Chain length determination
T303_06370	<i>epsJ</i>	Glycosyl transferase
T303_06375	<i>epsI</i>	Glycosyl transferase
T303_06380	<i>epsH</i>	Glycosyl transferase
T303_06385	<i>epsG</i>	Glycosyl transferase
T303_06395	<i>epsF</i>	Rhamnosyl transferase
T303_06400	<i>epsE</i>	Glycosyl transferase
T303_06405	<i>eps1D</i>	Chain length determination
T303_06410	<i>eps1C</i>	Chain length determination
T303_06415	<i>epsB</i>	Regulation
T303_06420	<i>epsA</i>	Regulation



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