

Figure 1

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Figure 2

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101 .....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
   DDSSFSRVQVP SHLETAGLLAPQYVNVQYPWDGHEDPKAPAIPEHGHVAVYRREFDADGEVAQAVREGRPVTLTFQGAATAIYVWLNGSFVGYAEDSFTP

      210     220     230     240     250     260     270     280     290     300
201 .....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
   SEFDVTDIAIKVDGNVLAVVCYESSASWLEDQDFWRLHGLFRSVELNARPAAHIALHADADNDLATSRGSLSLDLIDGAANAATVDFALWOKNGTIVW

      310     320     330     340     350     360     370     380     390     400
301 .....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
   HTATKADGTLHAEEIDDAAPWSAERPDLVELSVTLLDAGKVLETARTRIGFRHVAIEDGILKLNKRLVFRGVNRHFPDCRRGRAITEEDMLWDIRFM

      410     420     430     440     450     460     470     480     490     500
401 .....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
   KRHNINAVRTSHYPNQSRWYELCDEYGIYLIIDETNLETHGSWNSPGDIPVGTSPVGDEAWLGACIDRLDSMILDRNHPSVLVWSLGNESYAGEVLKAM

      510     520     530     540     550     560     570     580     590     600
501 .....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
   SAHAHRLDPGRFVHYEGVNNHAYDGISDFESRMYPKPAEIQDWLEHGDERGEASKPFVSCEYMHAMGNSCGGLSEFIDLERYERYSGGFIWDYIDQGLV

      610     620     630     640     650     660     670     680     690     700
601 .....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
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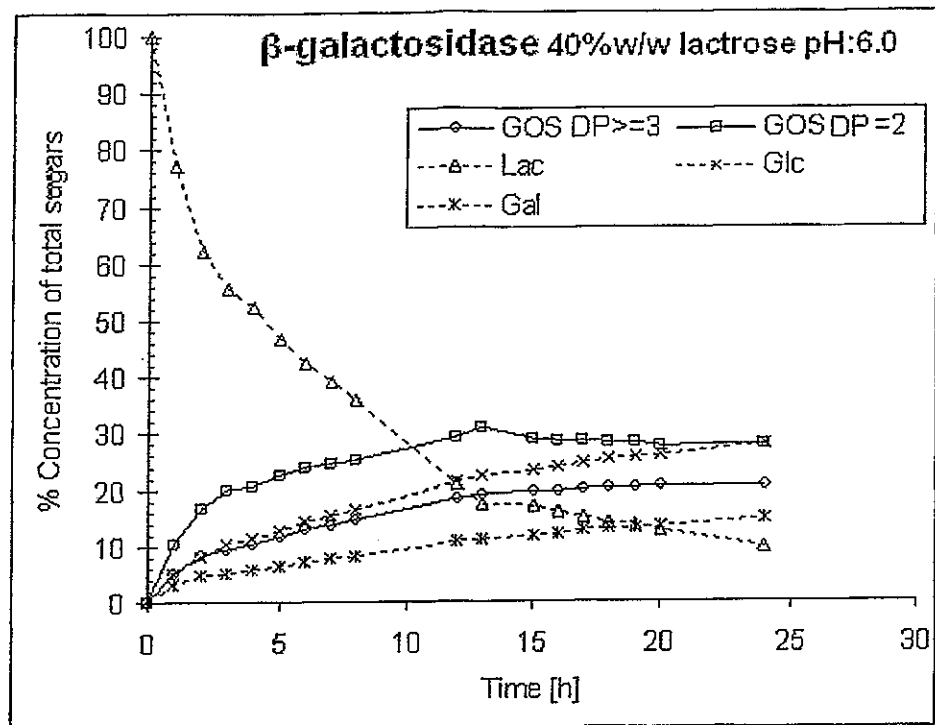
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701 .....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
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      810     820     830     840     850     860     870     880     890     900
801 .....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
   DRENVIRPELVTFRPLTDNDRGNHSGFDRAAWFAAGRYAIVTETKIHESDDGLVAEYQYELADPNHTPVSVTYHVNSDMRMQLTVEYPGNATDMASLPA

      910     920     930     940     950     960     970     980     990    1000
901 .....|.....|.....|.....|.....|.....|.....|.....|.....|.....|
   FGIWELPGEYDRLRYYPGPEETRYDRKQGGKLGIWDATAKASMAPYLMVQETGSHEDVRWLEATDIQGHGLRVTQRGDRHFTASLLPWNTYIEAARR

      1010    1020    1030    1040    1050
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5 Figure 3

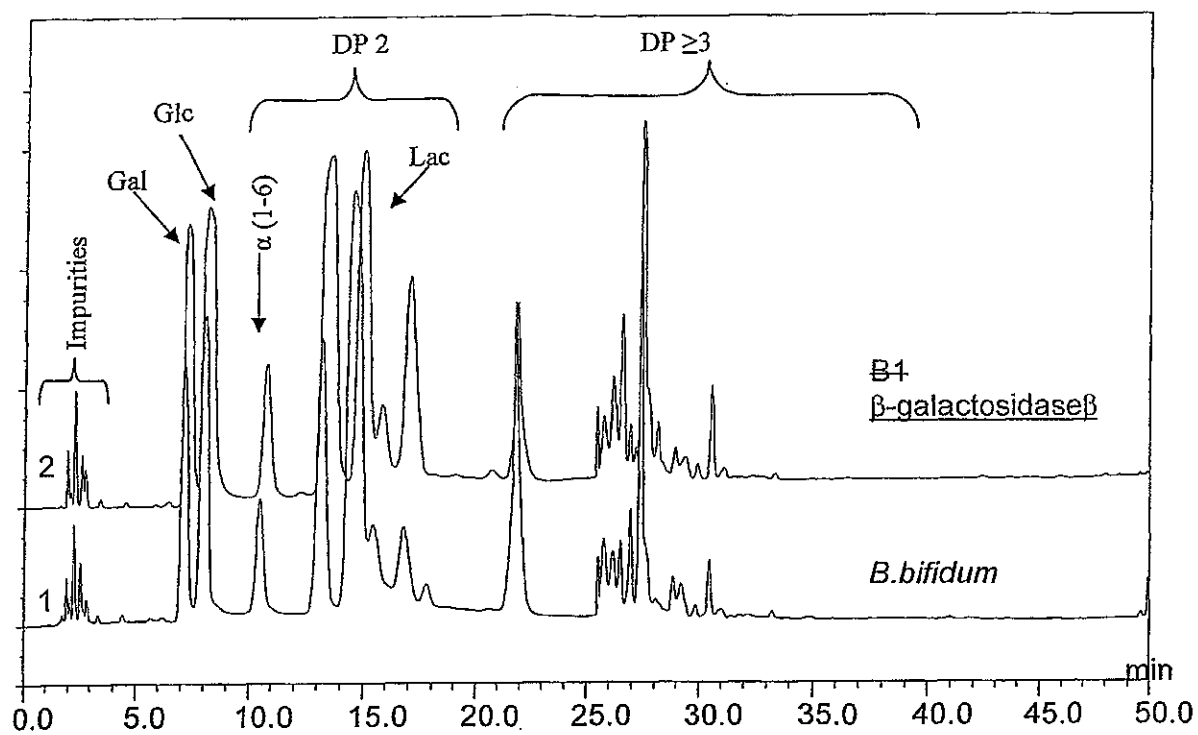


Figure 4