

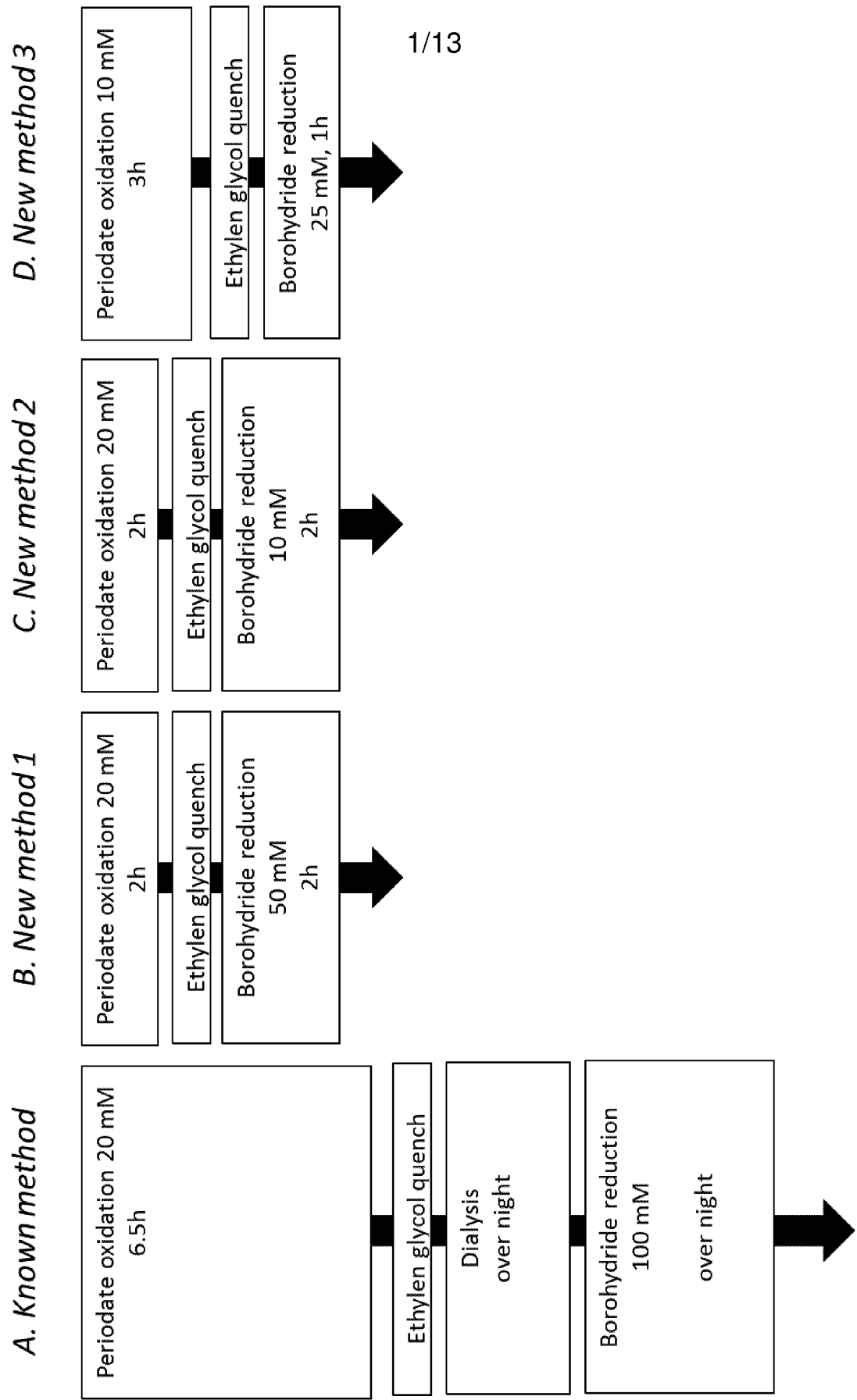


Figure 1

Known method *New method 1*

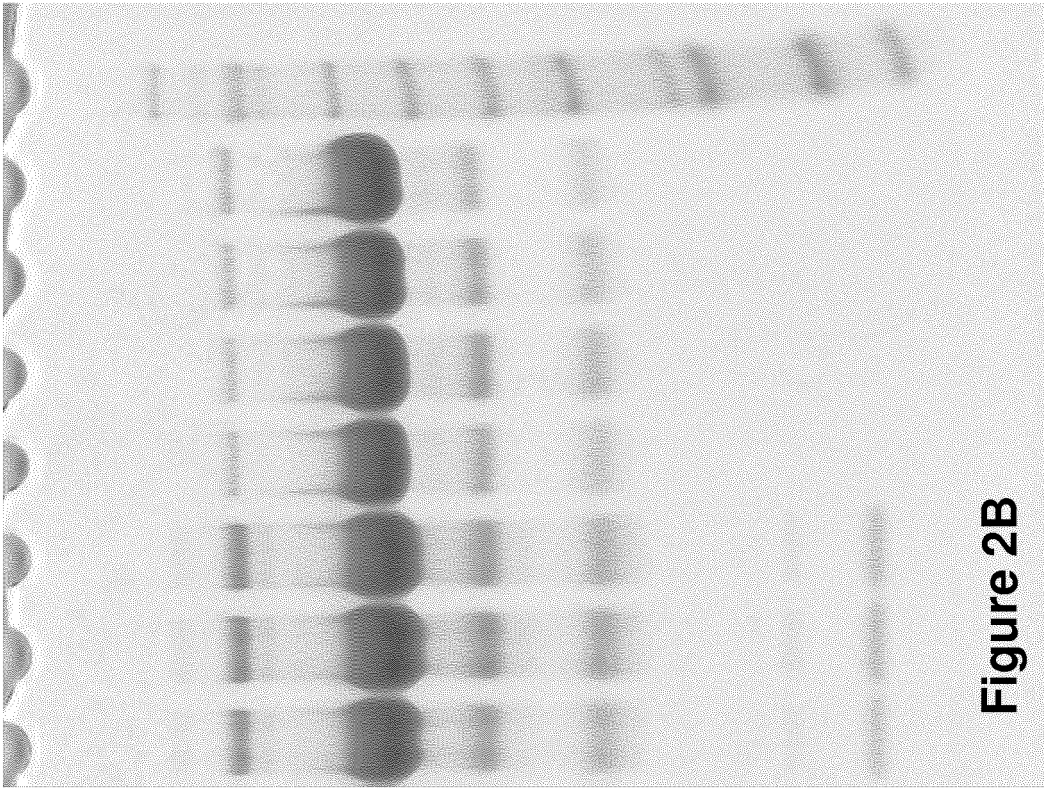


Figure 2B

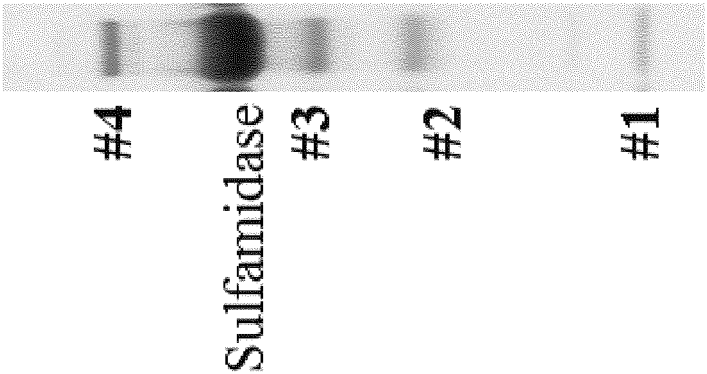
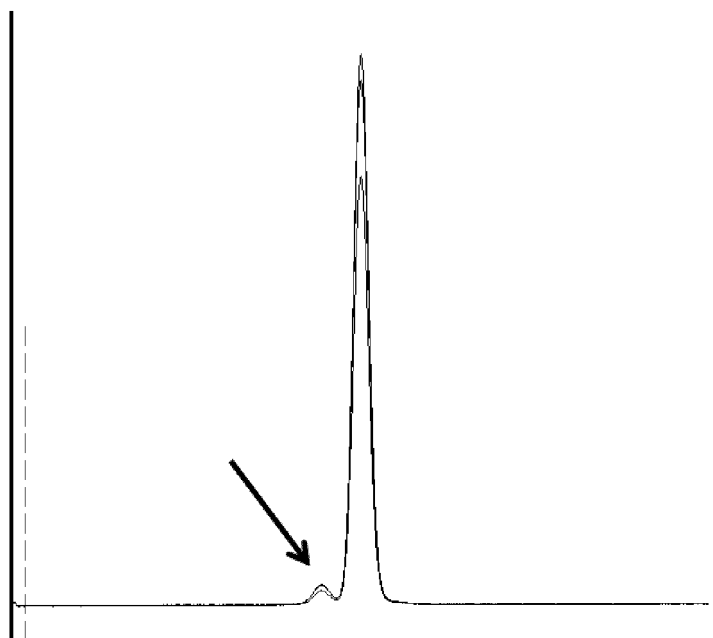


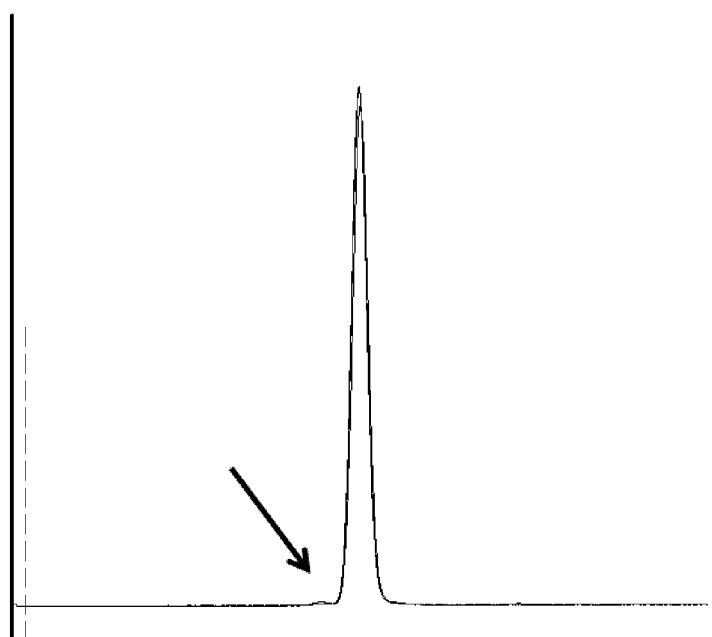
Figure 2A

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A

Known method

B

New method 1**Figure 3**

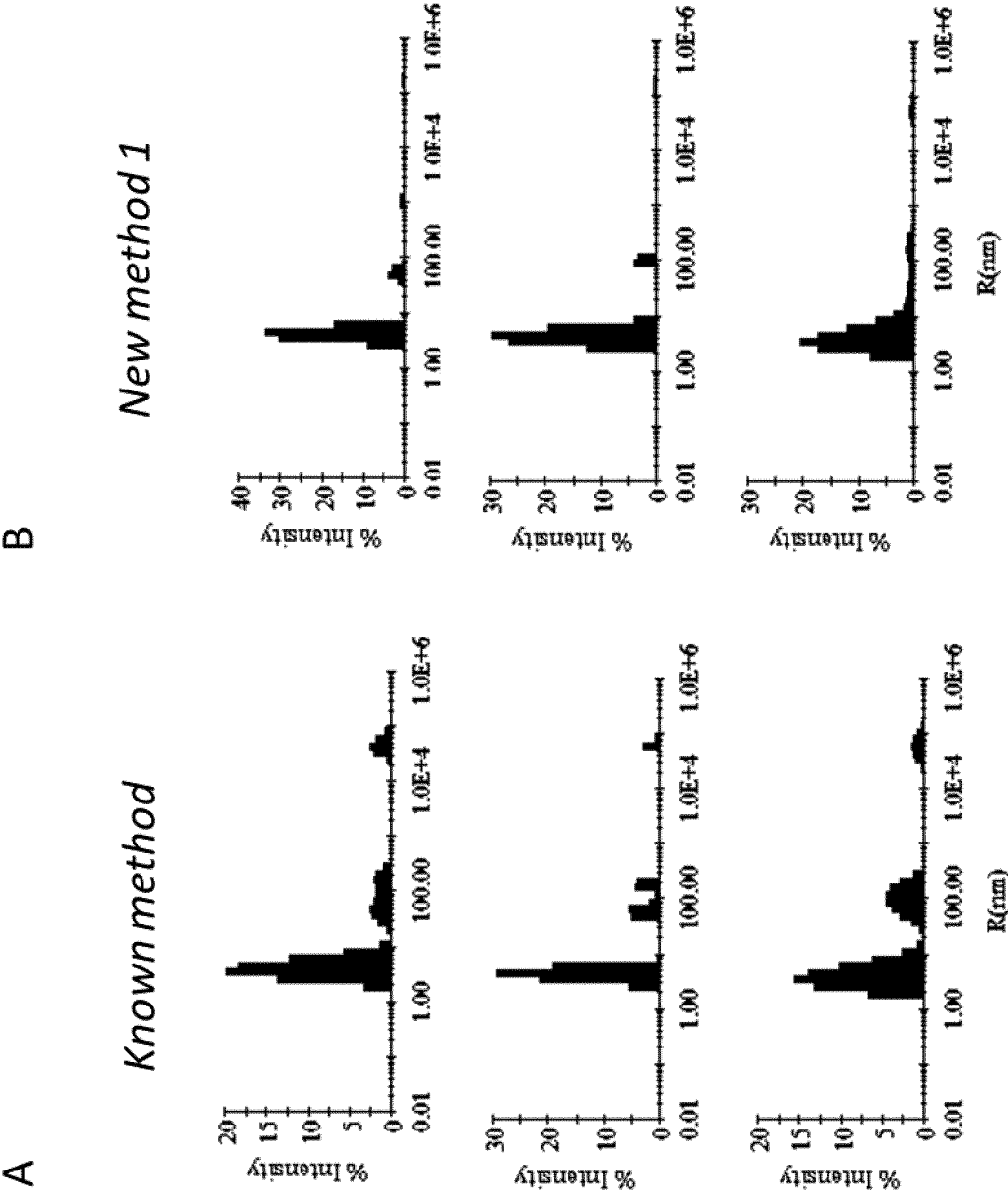


Figure 4

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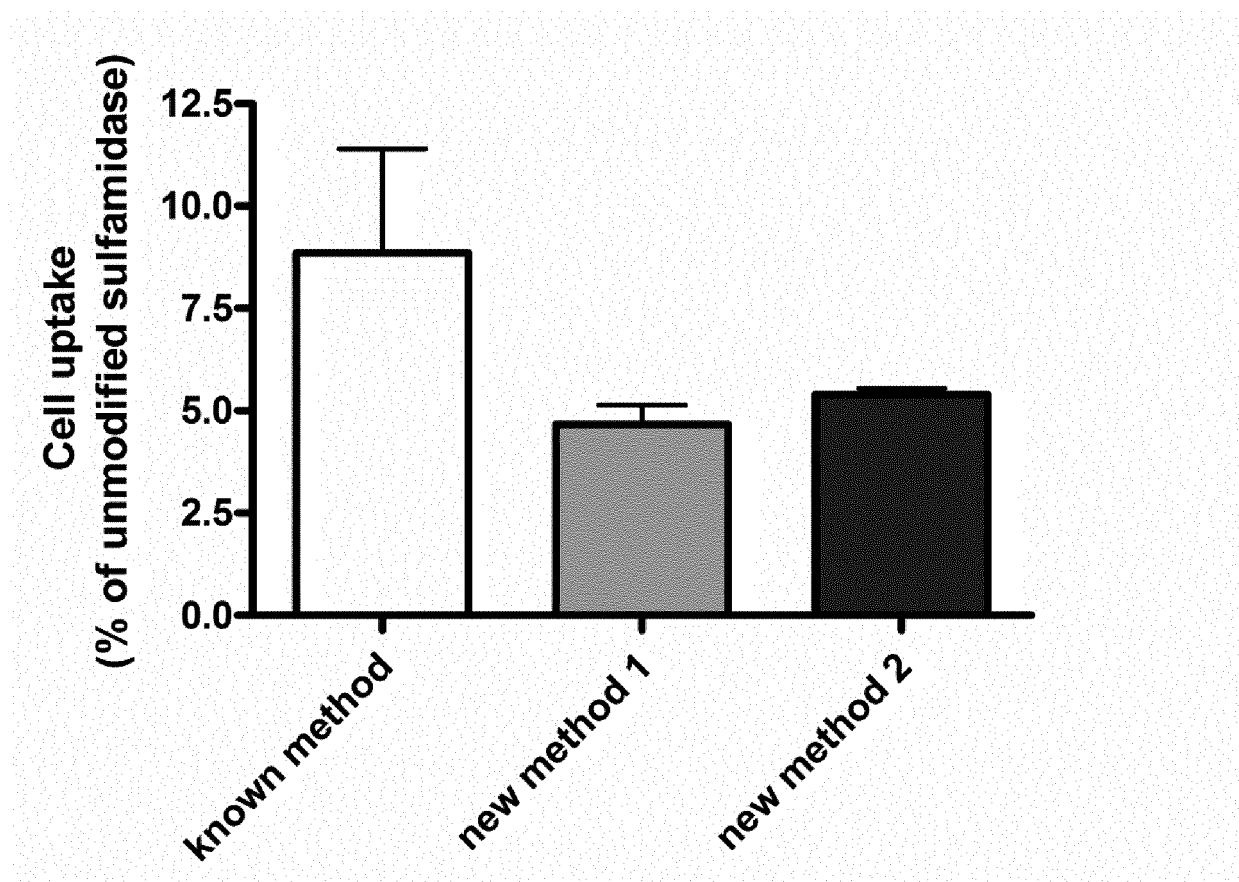
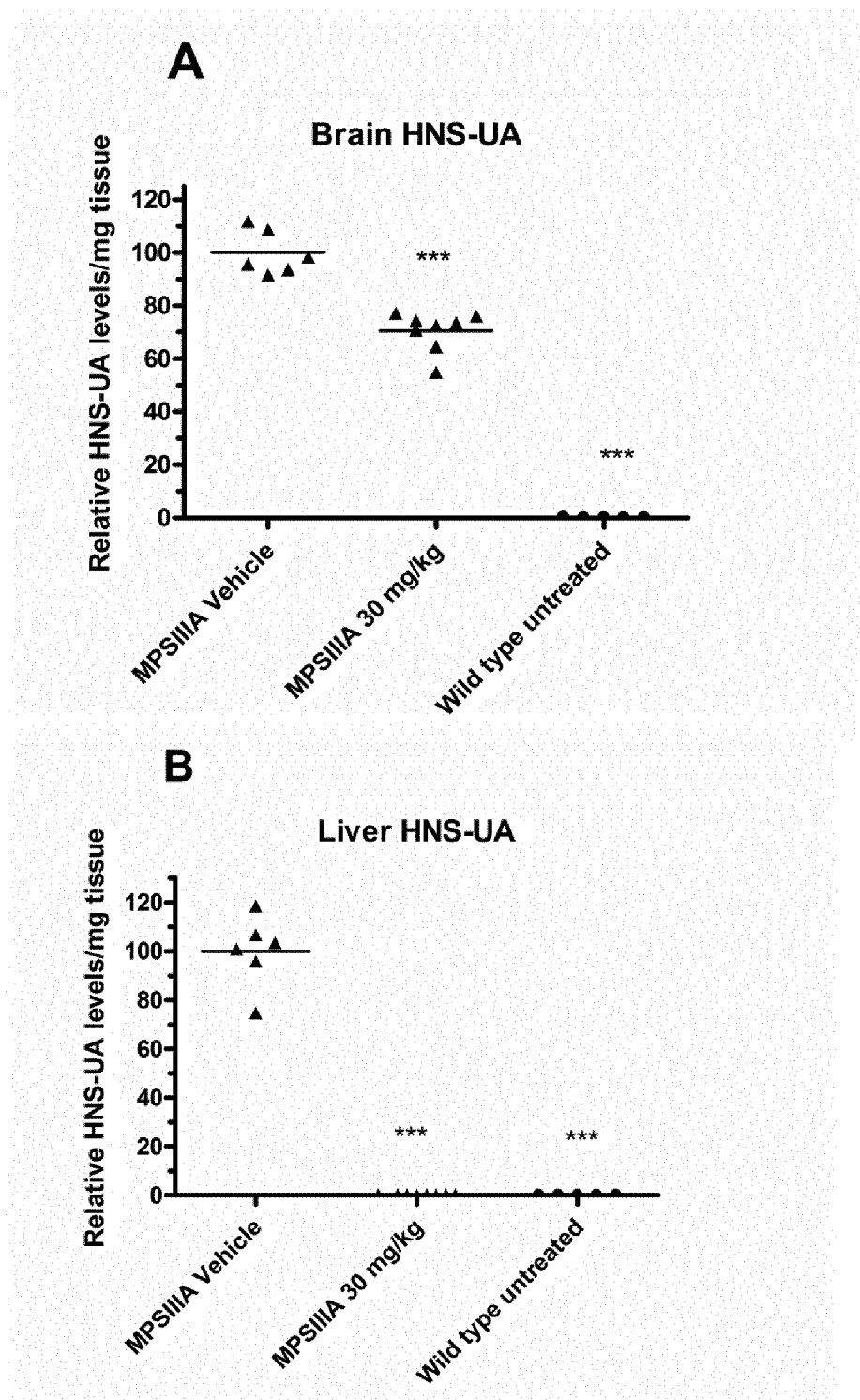


Figure 5

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**Figure 6**

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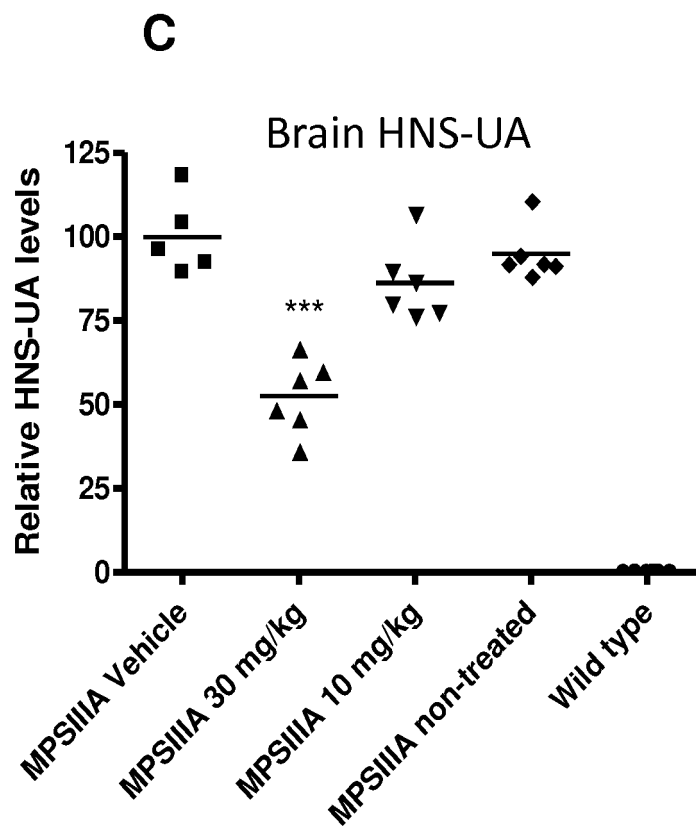


Figure 6

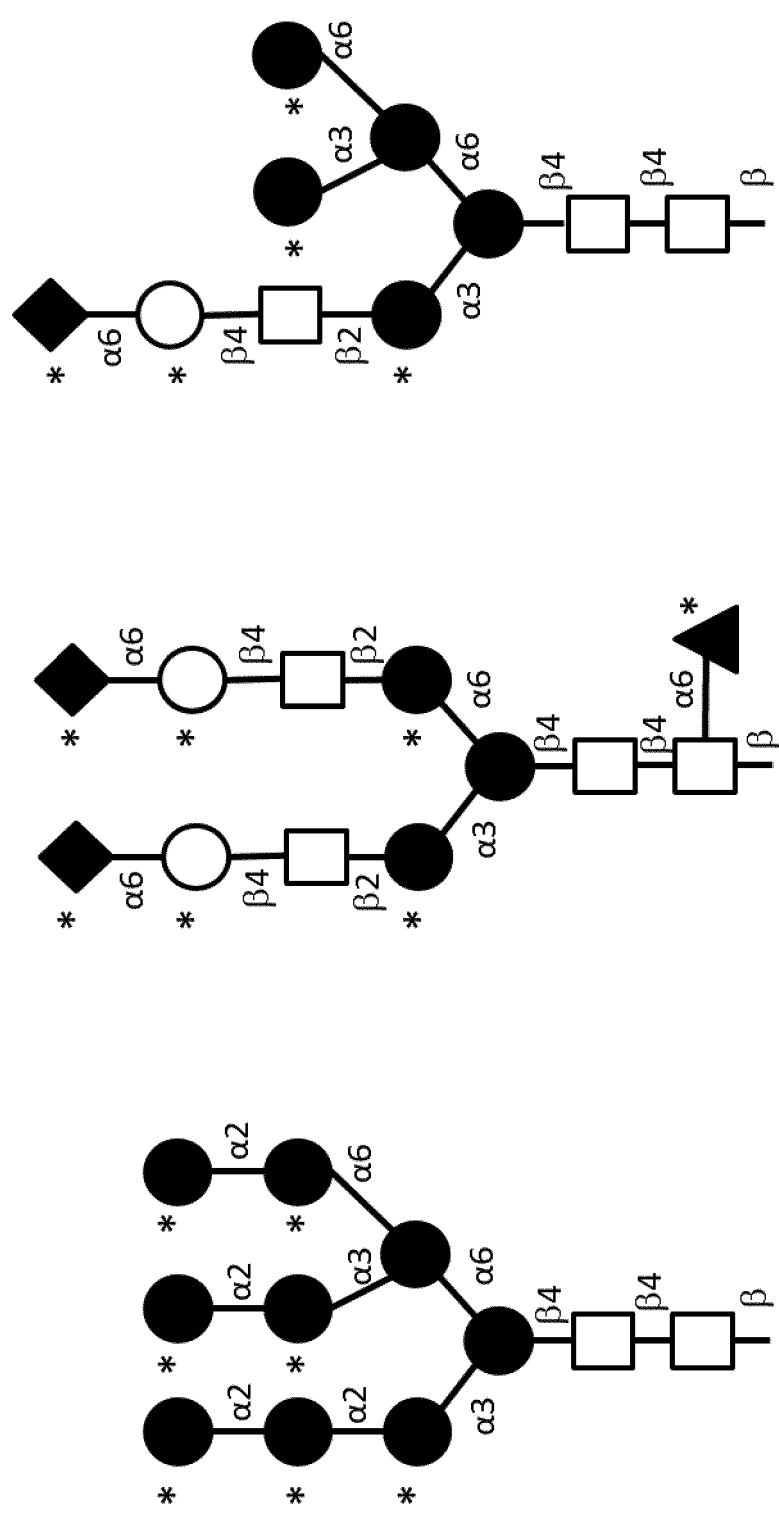


Figure 7

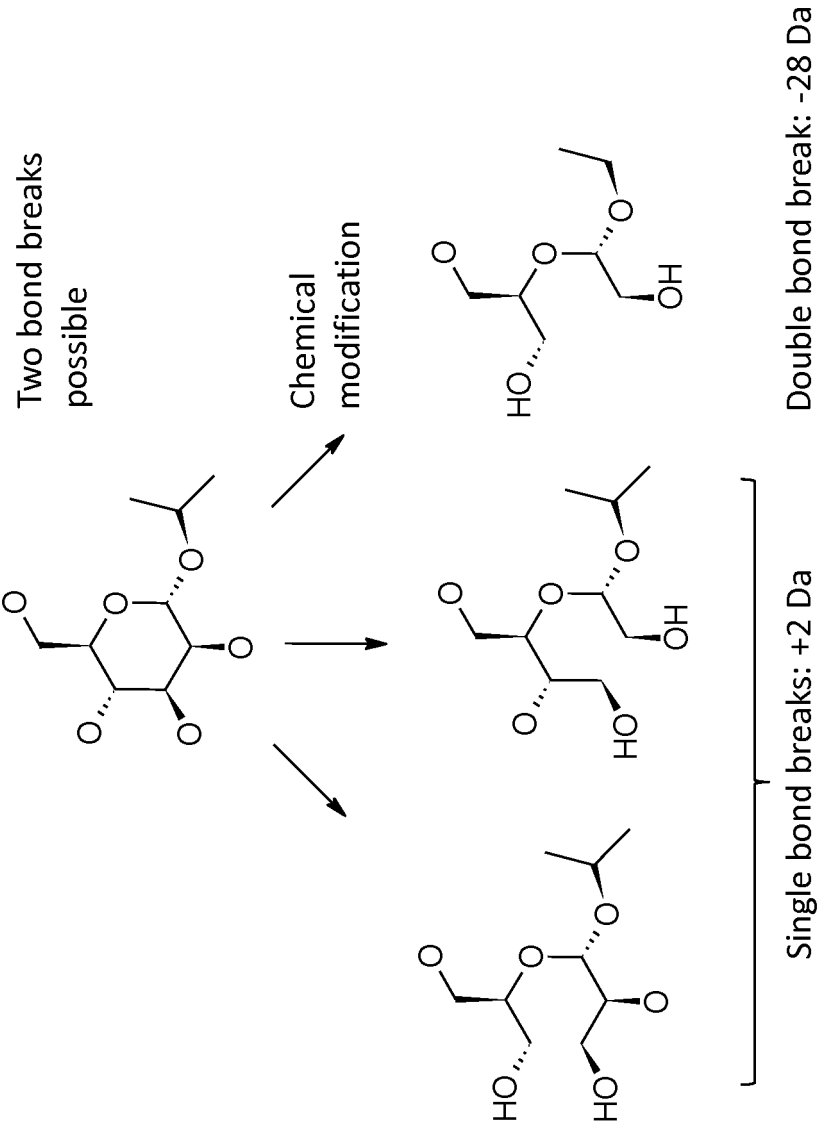


Figure 8A

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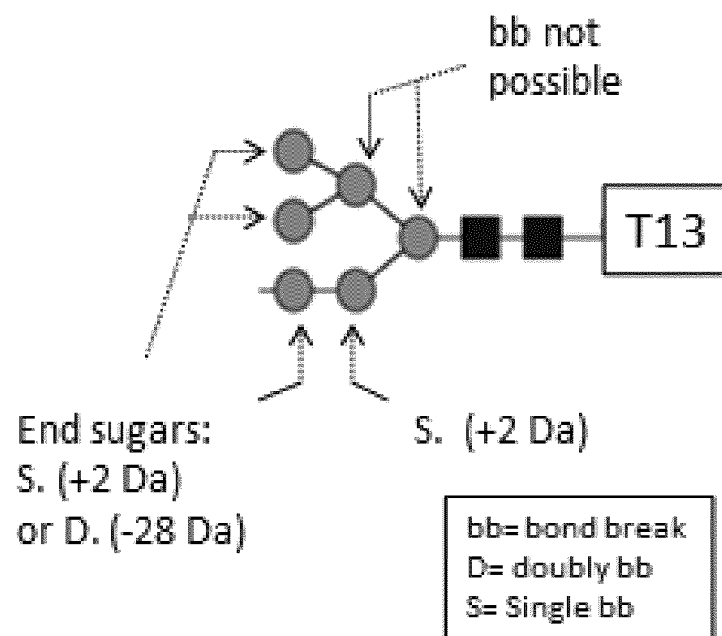


Figure 8B

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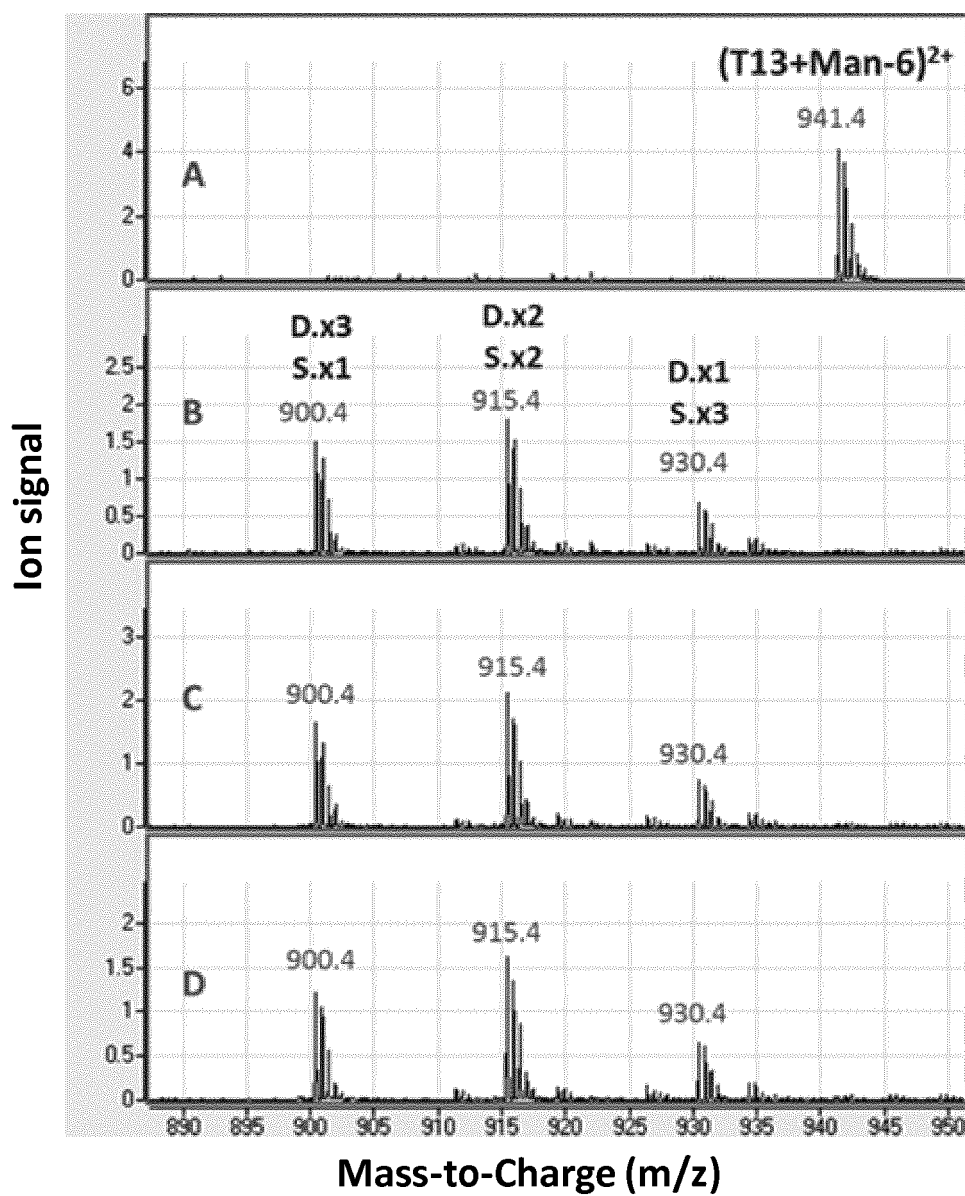
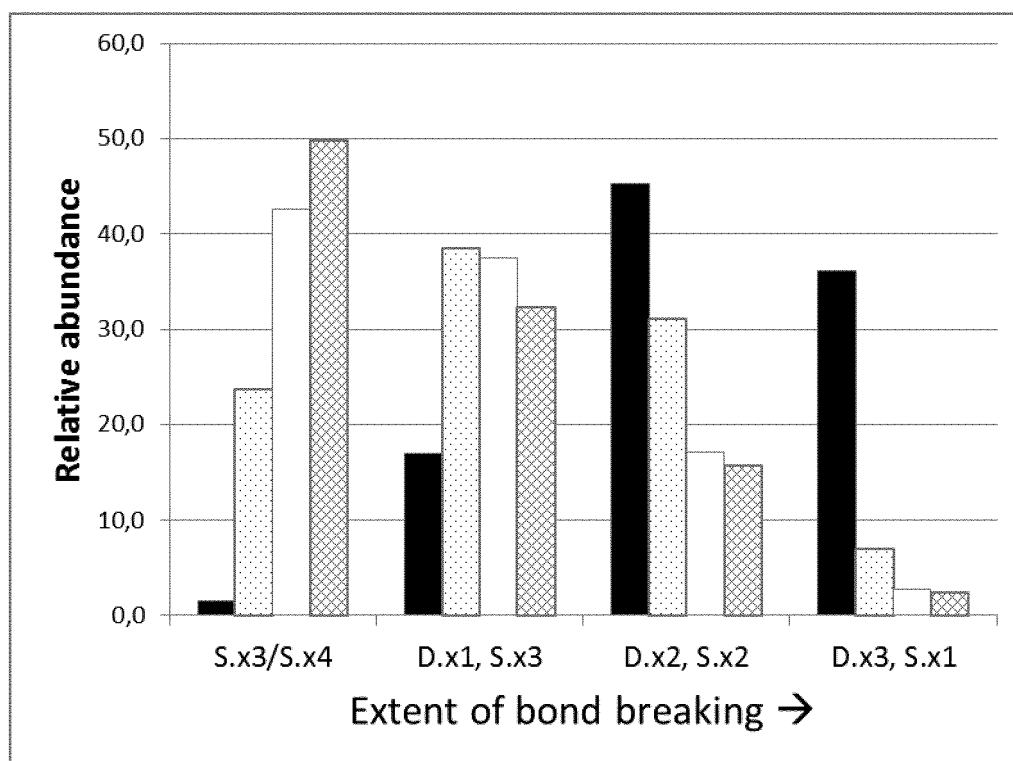
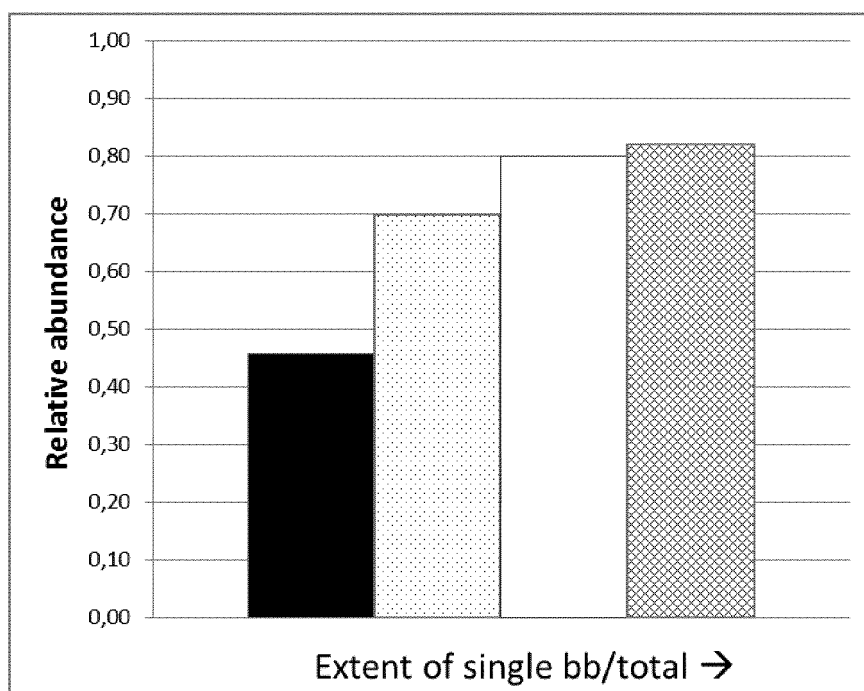


Figure 9

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**A****B****Figure 10**

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SEQ ID NO:	Amino acid sequence	Name
1	RPRNALLLADGGFESGAYNNSAIAIATPHLDALARRSLLFRNAFTSVSSCSPSRASLLTGLPQHONG MYGLHQDVHHFNSFDKVRSLPLLLSQAGVRTGIIGKKHVGPEVYVYFDFAYTEENGSVLQVGRNITRI KLLVRKFLQTQDDRPFFLYVAFHDPHRCGHSQPQYGTCEKFGNGESGMGRIPDWTTPQAYDPLDV LVPYFVPNTPAARADLAAQYTTVGRMDQGVGLVQLERDAGVLDNLVFTSDNGIPFSPGRTNLYW PGTAEP LLVSSPEHPKRWGVSEAYVSLDLTPTILDWFSIPYPSYAIFGSKTIHLTGRSLLPALEAEP LWATVFGSQSHHEVTMSYPMRSVQHRHFRLVHNLNFKMPFPIDQDFYVSPTFQD LLNRTTAGQPTG WYKDLRHHYYRARWELYDRSRDPHETQNLATDPRFAQLLEMLRDQLAKWQWETHDPWVCAPDGV LEEKLSPPCCQPLHNEL	PBV2351
2	GSRPRNALLLADGGFESGAYNNSAIAIATPHLDALARRSLLFRNAFTSVSSCSPSRASLLTGLPQHONG NGMYGLHQDVHHFNSFDKVRSLPLLLSQAGVRTGIIGKKHVGPEVYVYFDFAYTEENGSVLQVGRNI TRIKLLVRKFLQTQDDRPFFLYVAFHDPHRCGHSQPQYGTCEKFGNGESGMGRIPDWTTPQAYDPL DVLVPYFVPNTPAARADLAAQYTTVGRMDQGVGLVQLERDAGVLDNLVFTSDNGIPFSPGRTNL YWPGTAEPLLVSSPEHPKRWGVSEAYVSLDLTPTILDWFSIPYPSYAIFGSKTIHLTGRSLLPALEA EPLWATVFGSQSHHEVTMSYPMRSVQHRHFRLVHNLNFKMPFPIDQDFYVSPTFQD LLNRTTAGQP TGWYKDLRHHYYRARWELYDRSRDPHETQNLATDPRFAQLLEMLRDQLAKWQWETHDPWVCAPD GVLEEKLSPPCCQPLHNEL	pBV1968
3	RPRNALLLADGGFESGAYNNSAIAIATPHLDALARRSLLFRNAFTSVSSCSPSRASLLTGLPQHONG MYGLHQDVHHFNSFDKVRSLPLLLSQAGVRTGIIGKKHVGPEVYVYFDFAYTEENGSVLQVGRNITRI KLLVRKFLQTQDDRPFFLYVAFHDPHRCGHSQPQYGTCEKFGNGESGMGRIPDWTTPQAYDPLDV LVPYFVPNTPAARADLAAQYTTVGRMDQGVGLVQLERDAGVLDNLVFTSDNGIPFSPGRTNLYW PGTAEP LLVSSPEHPKRWGVSEAYVSLDLTPTILDWFSIPYPSYAIFGSKTIHLTGRSLLPALEAEP LWATVFGSQSHHEVTMSYPMRSVQHRHFRLVHNLNFKMPFPIDQDFYVSPTFQD LLNRTTAGQPTG WYKDLRHHYYRARWELYDRSRDPHETQNLATDPRFAQLLEMLRDQLAKWQWETHDPWVCAPDGV LEEKLSPPCCQPLHNELGS	pBV1803

Figure 11