Figure 1:

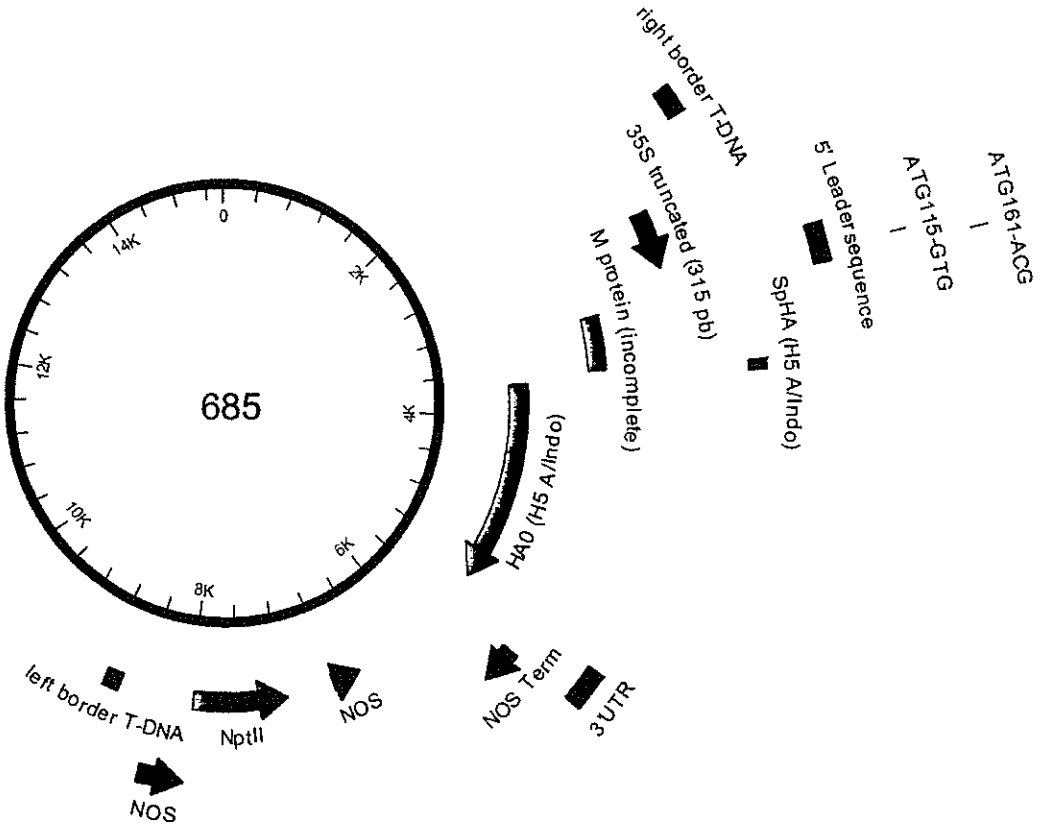

685
0
2K
4K
6K
8K
10K
12K
14K
left border T-DNA
NptII
NOS
NOS
HA0 (H5 A/Indo)
NOS Term
3'UTR
M protein (incomplete)
35S truncated (315 pb)
right border T-DNA
SpHA (H5 A/Indo)
5' Leader sequence
ATG115-GTG
ATG161-ACG

Figure 2A

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CGTGCTTCGGCACCAGTACAACGTTTTCTTTCACTGAAGCGAAATCAAAGATCTCTTTGTGGACACGTA
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CGCGCGGTGTCATCTATGTTACTAGATTCTAGAGTCTCAAGCTTCGGCGCGCC

Figure 2B

MEKIVLLLAIVSLVKSDQICIGYHANNSTEQVDTIMEKNVTVTHAQDILEKTHNGKLCDLDGVKPLILRDCS
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Figure 3A

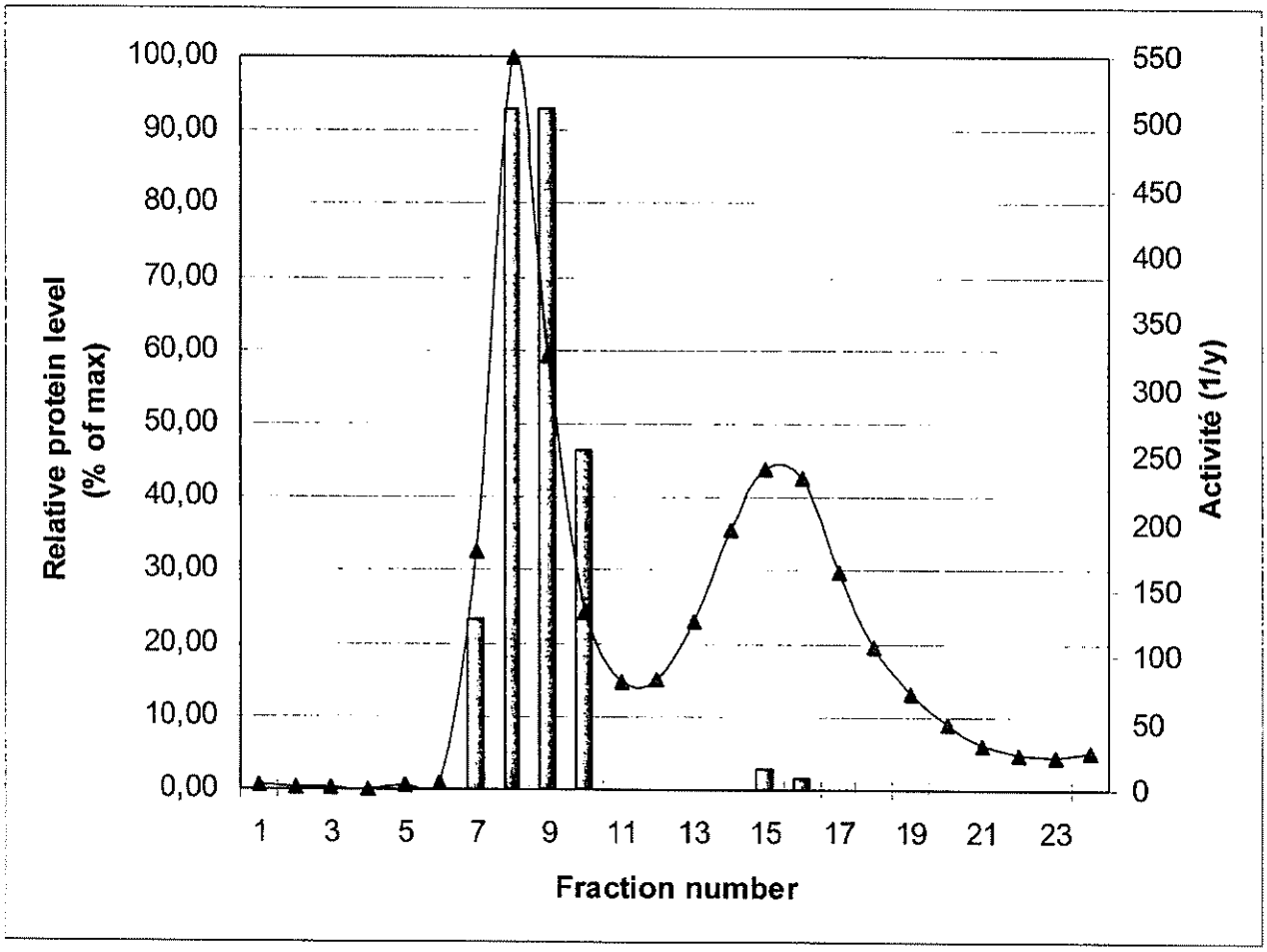

100,00
90,00
80,00
70,00
60,00
50,00
40,00
30,00
20,00
10,00
0,00
Relative protein level (% of max)
550
500
450
400
350
300
250
200
150
100
50
0
Activité (1/y)
1 3 5 7 9 11 13 15 17 19 21 23
Fraction number

Figure 3B

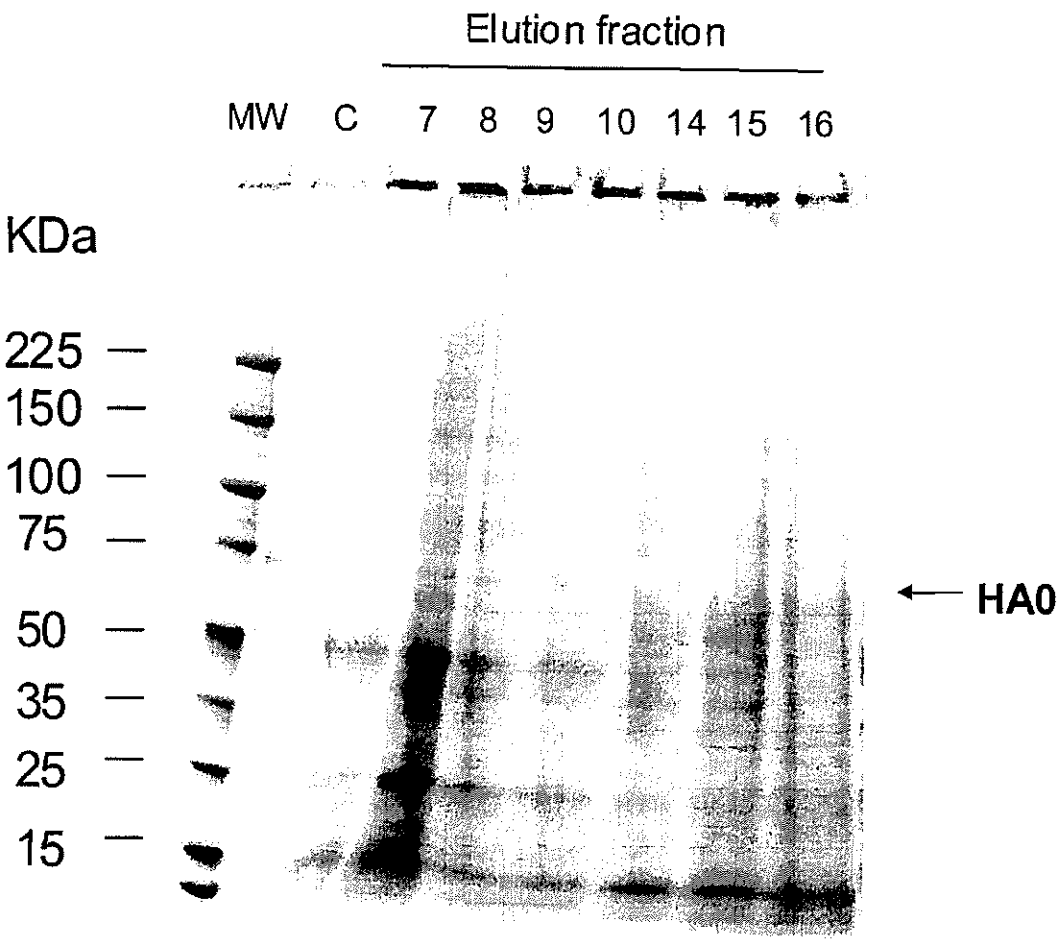

Elution fraction
MW
C
7
8
9
10
14
15
16
KDa
225
150
100
75
50
35
25
15
HA0

Figure 4

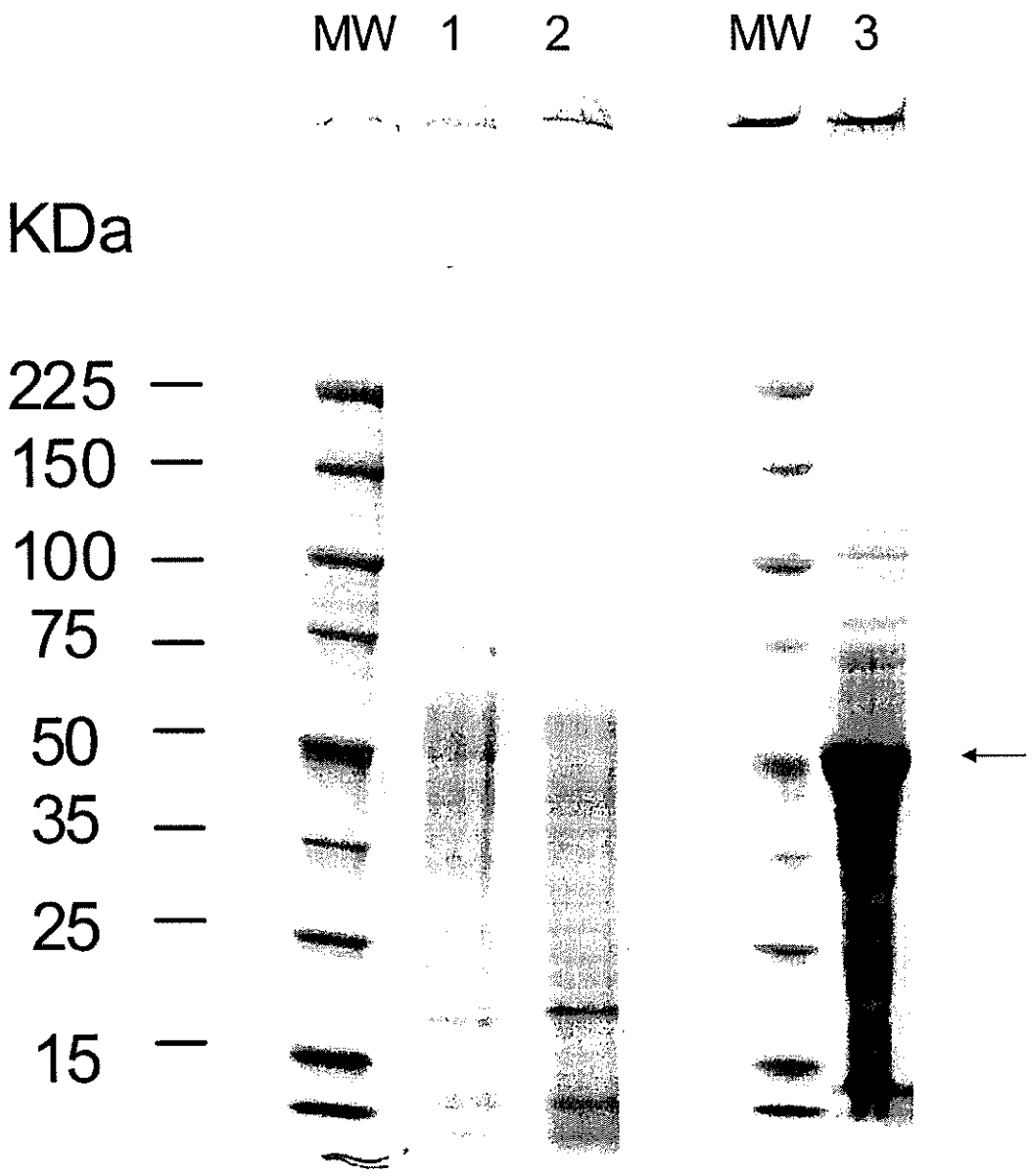
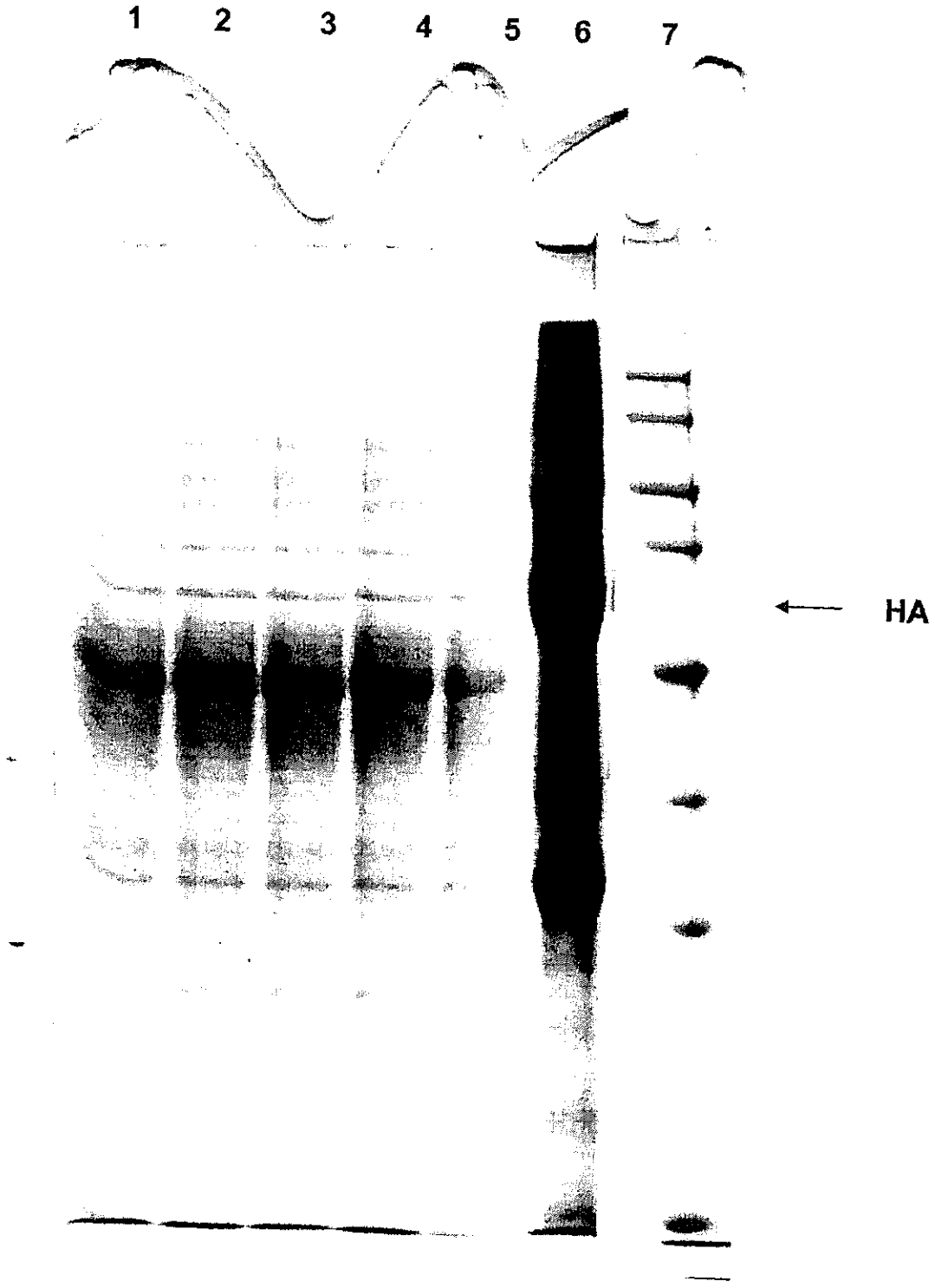


Figure 5

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Figure 6



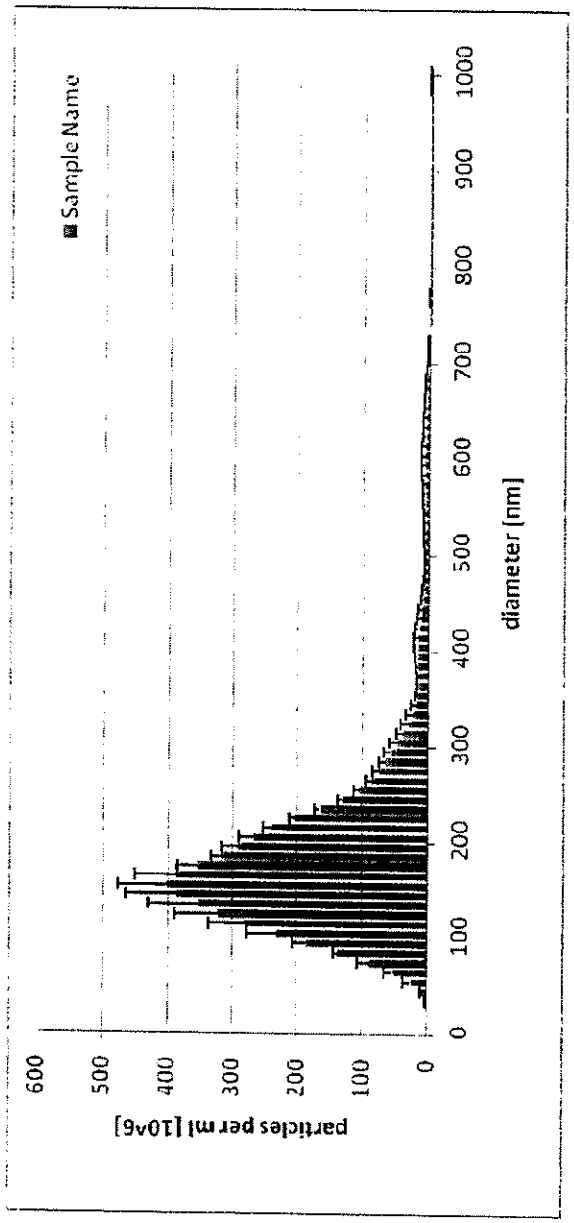

Sample Name
diameter [nm]
0
100
200
300
400
500
600
700
800
900
1000
particles per ml [10e6]
600
500
400
300
200
100
0

Figure 7A

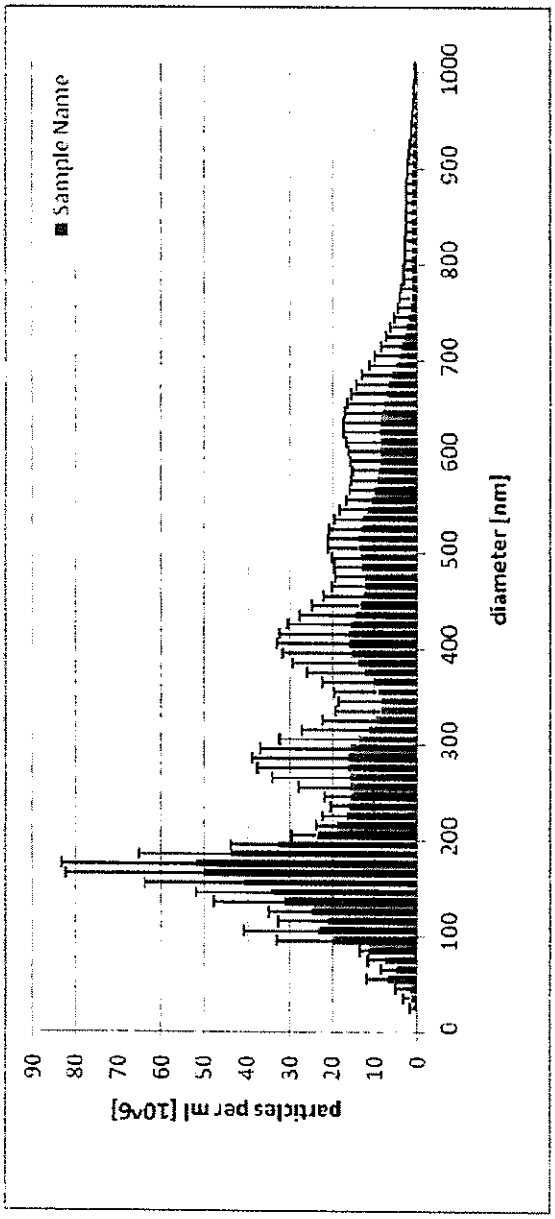

Sample Name
particles per ml [10^6]
diameter [nm]
0
10
20
30
40
50
60
70
80
90
0
100
200
300
400
500
600
700
800
900
1000

Figure 7B

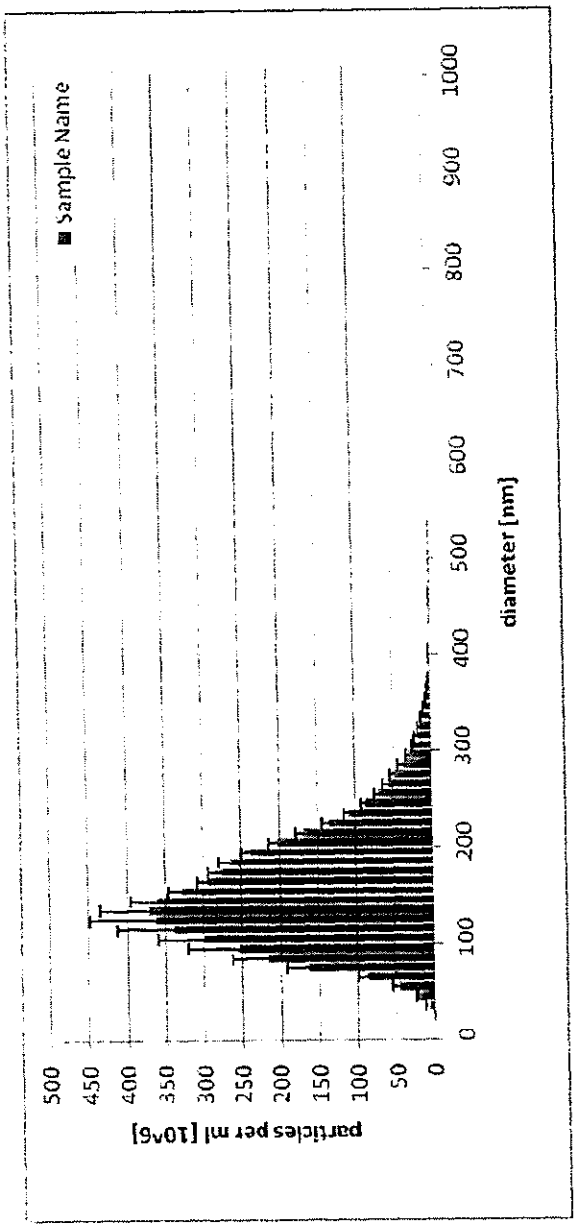

Sample Name
diameter [nm]
0
100
200
300
400
500
600
700
800
900
1000
particles per ml [10^6]
0
50
100
150
200
250
300
350
400
450
500

Figure 7C

Figure 8

