

<b>Supplementary Table 13</b>								
	<b>Number of binding sites on one transcript</b>							
<b>ID</b>	<b>2</b>	<b>3</b>	<b>4</b>	<b>5</b>	<b>6</b>	<b>7</b>	<b>8</b>	<b>9</b>
<b>6</b>	3	0	0	0	0	0	0	0
<b>13</b>	2	0	0	0	0	0	0	0
<b>21</b>	34	3	0	0	0	0	0	0
<b>31</b>	39	6	0	0	0	0	0	0
<b>32</b>	11	3	0	0	0	0	0	0
<b>35</b>	14	1	1	0	0	0	0	0
<b>40</b>	323	72	14	5	3	1	0	2
<b>50</b>	321	79	22	8	1	0	0	0
<b>52</b>	353	89	21	7	1	0	0	0

Supplementary Table 4 specifies the number of genes that show a sequence homology to the specific ASO at several positions within the sequence. In the column header the number of hits per gene is indicated.