

<b>Gene</b>	<b>Coordinates in genome (chromosome: start – stop, <u>according to</u> NCBI)</b>	<b>GenBank/DDBJ/EMBL Accession numbers</b>
<i>APOA1</i>	chr11:116835751 - 116837622	NG_012021.1
<i>APOA2</i>	chr1:161222293 - 61223628	NG_012043.1
<i>APP</i>	chr21:25880550 - 26171128	NG_007376.1
<i>B2M</i>	chr15:44711487 - 44718159	NG_012920.1
<i>CST3</i>	chr20:23627897 - 23638048	NG_012887.2
<i>FGA</i>	chr4:154583128 - 154590745	NG_008832.1
<i>GSN</i>	chr9:121268102 - 121332844	NG_012872.1
<i>ITM2B</i>	chr13:48233138 - 48262096	NG_013069.1
<i>LYZ</i>	chr12:69348354 - 69354233	NG_008195.1
<i>PRNP</i>	chr20:4686151 - 4701588	NG_009087.1
<i>TTR</i>	chr18:31591767 - 31599024	NG_009490.1

**Supplementary table 2:** List of genes included in the design together with their genomic coordinates and gene accession numbers