

## Familial Hypercholesterolaemia (R134) Genetic Testing Request Form

Patient Details			
<b>NHS No:</b>		<b>Sex:</b>	
<b>Surname:</b>		<b>Address:</b>	
<b>Forename:</b>			
<b>Date of Birth:</b>		<b>Postcode:</b>	
<b>Hospital:</b>		<b>Hospital No:</b>	
<b>Ethnicity:</b>			

Requester Details			
<b>Clinician:</b>		<b>Email:</b>	
<b>Reporting Address:</b>		<b>Invoice Address:</b>	

Clinical Information					
CVD History	Age	Clinical signs of FH	Lipids		
ACS/MI		Corneal arcus		<b>Currently on statins?</b>	
CABG		Xanthelasma		<b>Dose:</b>	
PTCA		Tendon xanthoma		<b>Pretreatment levels:</b>	
Angina				<b>Total cholesterol:</b>	
Stroke/TIA				<b>LDL-c</b>	
PVD				<b>HDL-c</b>	
<b>Carotid artery intima-media thickness:</b>				<b>Triglycerides</b>	
<b>Other:</b>					
<b>Referral criteria used:</b>	Simon Broome:	Welsh:	Dutch:	Other:	<b>Score:</b>
<b>Family history of CVD, raised cholesterol etc.</b>					

Testing required					
<b>Diagnostic testing</b> <ul style="list-style-type: none"> <li>Full sequence analysis of LDLR, APOB, PCSK9, LDLRAP1, APOE</li> <li>Dosage analysis of LDLR</li> <li>Polygenic LDL-C-raising SNP score</li> </ul> Please note dosage analysis may be less reliable on DNA from buccal swabs. Blood is the preferred tissue type for this analysis.					
<b>Familial variant testing</b> Testing for a known familial variant for either: <ul style="list-style-type: none"> <li>Cascade testing of a pathogenic variant</li> <li>Segregation analysis of a variant of uncertain significance</li> </ul>	<table style="width: 100%; border-collapse: collapse;"> <tr> <td style="width: 50%; border-bottom: 1px dashed black;">Affected:</td> <td style="width: 50%; border-bottom: 1px dashed black;">Unaffected:</td> </tr> <tr> <td colspan="2" style="border-bottom: 1px dashed black;">Affected patients only</td> </tr> </table>	Affected:	Unaffected:	Affected patients only	
Affected:	Unaffected:				
Affected patients only					
Index case name:					
Index case date of birth:					
Relationship to the patient:					
Please provide a copy of the relative's diagnostic genetic report or as much information as possible regarding where and when testing was carried out including the variant if known					