Elucigene FH20 and LIPOchip for FH - data extraction form

Reviewer ID: Date: Administration details Study ID: Publication status: Other papers this study may link with: Aim of the study Test(s) reported **Cascade testing Index cases** Elucigene LIPOchip LDL-C Targeted gene sequencing *CGA * includes DNA sequence analysis+ test for deletion/duplication+ analysis of APOB p.Arg3527Gln and PCSK9 p.Asp374Tyr using various techniques. Outcomes reported Diagnostic accuracy Mutation detection rate Clinical effectiveness Study details Cross-sectional comparative RCT Case control study Cross-sectional single test Other, please specify: Multicentre study? Yes No If Yes, number of centres: Consecutive recruitment? Yes No Not stated Country: Study dates: Length of follow up: Source of funding: Inclusion criteria:

Exclusion criteria:			
Baseline characteristics of	^c participants		
Adults Chi	ldren		
Criteria used for clinical di	agnosis:		
Simon Broome			
Dutch			
Medped			
Other	specify the LDLC cu	t offs used and definition	
Type of FH:			
Possible	Unclassified F	Н	
Definite	Not stated		
Heterozygous FH	Homozygous	FH	
Diagnosis of the Index case	es confirmed by:		
Clinical test	Genetic test		
Who perform the clinical d	iagnosis?		
Number of participant/san	nple		
give detail of each type of FH if reported	All	Index cases	Relatives
Eligible			

Enrolled			
Analysed			
Received index test(s)			
Received comparator test(s) for index cases			
Received cascade test(s) 1 st degree relatives 2 nd degree relatives 3 rd degree relatives			
Received comparator test(s) 1 st degree relatives 2 nd degree relatives 3 rd degree relatives			
Age (mean/ median, SD, range)			
Receiving treatment for hyper- cholestorelaemia (specify treatment)			
Ethnicity			
Gender	M: F:	M: F:	M: F:
Tendon xanthomas			
Coronary Heart Disease			
Intervention tests Elucigene FH20 (Tepnel	molecular diagnosti	cs)	
If not FH20 which version	and how many mutat	tions was it designed to detect?	
Gel-based analysis	Fluorescent anal	ysis:	
Who carried out the test?			
Where was the test underta	ıken?		

Time taken to obtain test results:
Additional information on the test:
LIPOchip (Progenika Biopharma)
If not version 10 which version and how many mutations was it designed to detect?
Samples processed at: LIPOchip laboratory Other If other, please give details:
Methodology used:
DNA array
Analysis for large gene re-arrangements
Automated sequencing of the LDLR
Who carried out the test?
Where was the test undertaken?
Time taken to obtain test results:
Additional information on the test:
Comparator tests
CGA (as defined on page 5 of the protocol)
CGA should include following:
DNA sequence analysis of the promoter, all exons, the exon/intron boundaries and into 3' untranslated region of the LDLR gene
Manufacturer and any other technical characteristics of the test:
MLPA for each exon and the promoter region of the LDLR gene to detect deletions and duplications
Manufacturer and any other technical characteristics of the test:

Analysis for the common APOB p.Arg3527Gin and PCSK9 p.Asp374Tyr gene mutations
Manufacturer and any other technical characteristics of the test:
Who carried out the test?
Where was the test undertaken?
Time taken to obtain test results:
Additional information on the test:
LDL-C concentration
LDL-C concentration Estimated from a fasting blood sample using the Friedwald equation? Yes No
LDL-C concentration Estimated from a fasting blood sample using the Friedwald equation? Yes No If No please specify method used:
Estimated from a fasting blood sample using the Friedwald equation? Yes No
Estimated from a fasting blood sample using the Friedwald equation? Yes No
Estimated from a fasting blood sample using the Friedwald equation? Yes No If No please specify method used:
Estimated from a fasting blood sample using the Friedwald equation? Yes No If No please specify method used: For cascade test please specify age and gender specific LDL-C cut offs:
Estimated from a fasting blood sample using the Friedwald equation? Yes No If No please specify method used: For cascade test please specify age and gender specific LDL-C cut offs: No. of times LDL-C was measured? Once Twice Not stated
Estimated from a fasting blood sample using the Friedwald equation? Yes No If No please specify method used: For cascade test please specify age and gender specific LDL-C cut offs: No. of times LDL-C was measured? Once Twice Not stated Criteria used to define a positive test result:

	ation on the	test:				
If targeted gene se	equencing o	f relatives	was undert	aken, please	give details:	
				_		
Reference standar	d tast					
Kejerence standart	u test					
Was there a referen	ice standard	test that co	nsisted of e	ither of the fo	llowings?	
CGA in combination	on with Sim	on Broome	criteria		7	
CGA in combination	on with only	on broome	Citteria		_	
CGA only						
Simon Broome onl	V]	
					_1	
Results for Index	cases					
Genetic te	st					
	LDLR	APOB	PCSK9	MLPA	sequencing	Total
Number of						
Number of participants						
participants Number of						
participants						
Number of samples analysed						
participants Number of						
Number of samples analysed n/N (%) with mutation detected						
n/N (%) with						
n/N (%) with mutation detected						
participants Number of samples analysed n/N (%) with mutation detected n/N (%) with no mutation detected	est					
participants Number of samples analysed n/N (%) with mutation detected n/N (%) with no mutation detected 2. Genetic to Number of	est					
participants Number of samples analysed n/N (%) with mutation detected n/N (%) with no mutation detected 2. Genetic to	est					
participants Number of samples analysed n/N (%) with mutation detected n/N (%) with no mutation detected 2. Genetic to Number of	est					
participants Number of samples analysed n/N (%) with mutation detected n/N (%) with no mutation detected 2. Genetic to Number of participants	est					
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participants Number of samples analysed n/N (%) with mutation detected n/N (%) with no mutation detected 2. Genetic to Number of participants	est					
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participants Number of samples analysed n/N (%) with mutation detected n/N (%) with no mutation detected 2. Genetic to Number of participants Number of samples analysed n/N (%) with mutation detected	est					
participants Number of samples analysed n/N (%) with mutation detected n/N (%) with no mutation detected 2. Genetic to Number of participants Number of samples analysed n/N (%) with	est					

3. LDL-C as	a clinical te	st				
Number of						
participants						
partito						
Number of						
samples analysed						
Number of FH						
diagnosed						
Number of FH not						
diagnosed						
diagnosed						
D 1/ C C 1						
Results for Cascad	e test					
Specify the genetic	test		•••			
Number of						
participants						
(Index cases)						
Number of						
samples analysed						
samples analysed						
N. 1 C						
Number of						
families tested						
n/N (%) with						
mutation detected						
n/N (%) with no						
mutation detected						
LDL-C age and sex	specific tos	+				
Number of	specific tes	l .				T
participants						
(Index cases)						
Number of						
samples analysed						
Number of						
families tested						
rammes tested						
Number of FH						
diagnosed						
Number of FH not						
diagnosed						
Record data on each	level of an	alysis conta	aining 2x2 t	ables of true	and false positive	es and negatives
for					1	Č
101						

Test accuracy of genetic test (Elucigene/Lipochip) vs genetic test (CGA)

Test accuracy of genetic test (Elucigene/Lipochip) vs clinical test (LDL-C-SB criteria)

Subgroup analysis reported (e.g. ethnicity)? Yes No
If Yes please give details:
Give details of any clinical effectiveness outcomes reported, e.g. cholesterol levels following
treatment, CHD events etc or probability of true FH:
tremment, care events the or prosummy or true and
Give details of any information reported on acceptability and/or interpretability of the tests
or the details of any information reported on acceptability and/or interpretability of the tests
Additional information:
Audiuonai miofinauon;