

Viapath Analytics Molecular Pathology Laboratory

Please send all samples to: Molecular Pathology c/o Central Specimen
 Reception, Blood Sciences Laboratory, Ground Floor Bessemer Wing, King's College
 Hospital, Denmark Hill, London SE5 9RS

Molecular Pathology Referral Form

Patient details		Referrer details	
Surname		Surname	
First name		First name	
Date of Birth		Department	
Sex		Address line 1	
Ethnicity		Address line 2	
NHS number		Address line 3	
Your Hospital no		Postcode	
Your Lab no		Telephone	
Antenatal (<i>ANT</i>)	Yes <input type="checkbox"/> No <input type="checkbox"/>	Email	
Partner (<i>if ANT</i>)			

Pathology Results (please provide as much detail as possible).

Iron/liver Parameters		Haematology indices		
Serum Ferritin		Hb		HbF %
Serum Iron		RBC		HbA ₂ %
Serum TSat		MCV		Hb variant %
Serum Bilirubin		MCH		Absolute Reticulocyte
		Platelets		Reticulocyte %
Blood Film comments:				
Reason for referral/ family details				

Sample requirements:

For haemoglobinopathy investigation:	2 x 4 ml EDTA blood
Children and adults (all other tests):	4 ml EDTA blood
Infants:	1 ml EDTA blood
As DNA for Next Generation Sequencing:	3-5µg genomic DNA
As DNA for all other tests:	1-5µg genomic DNA

Date sample collected:

Molecular Tests (please tick all that apply)

- | | |
|---|---|
| <input type="checkbox"/> Hb variant identification | <input type="checkbox"/> Gilbert's genotyping (TA _{5/6/7/8} repeat) |
| <input type="checkbox"/> Haemoglobinopathy investigations | <input type="checkbox"/> Hereditary haemochromatosis (HFE)
(C282Y and H63D variants) |
| <input type="checkbox"/> Alpha thalassaemia | <input type="checkbox"/> Alpha-1-antitrypsin genotype
(S and Z alleles) |
| <input type="checkbox"/> Beta thalassaemia | <input type="checkbox"/> Thrombophilia genetic screen (please tick
all that apply):
FVL <input type="checkbox"/> PT <input type="checkbox"/> MTHFR <input type="checkbox"/> |
| <input type="checkbox"/> Pyruvate kinase gene sequencing | <input type="checkbox"/> Familial variant testing - for previously
identified variants from NGS (please state
familial variant): _____ |
| <input type="checkbox"/> Porphyria single gene sequencing
(please state gene): _____ | |
| <input type="checkbox"/> Other (please state): _____ | |

For further details of each test please refer to the [Viapath website](#)

Next Generation Sequencing

Red Cell Gene Panel

Please ensure FBC and film data are entered for all NGS requests.

Subpanels (please tick all that apply)

- | | |
|---|---|
| <input type="checkbox"/> Megaloblastic anaemia | <input type="checkbox"/> Sideroblastic anaemia |
| <input type="checkbox"/> Congenital dyserythropoietic anaemia | <input type="checkbox"/> Diamond-Blackfan anaemia |
| <input type="checkbox"/> Congenital erythrocytosis | <input type="checkbox"/> Haemoglobinopathies |
| <input type="checkbox"/> Red Cell Membranopathy | <input type="checkbox"/> Red Cell Enzyme |
| <input type="checkbox"/> Bone Marrow Failure | <input type="checkbox"/> Iron regulation |
| <input type="checkbox"/> Neutropenia | <input type="checkbox"/> HLH |
| <input type="checkbox"/> Thrombocytopenia | <input type="checkbox"/> Porphyria |
| <input type="checkbox"/> Lymphedema | |
| <input type="checkbox"/> Single gene analysis: _____ (name of gene) | |

For details of genes in each subpanel, please refer to the [Viapath website](#).

Patient consent

For all samples sent please ensure that the patient has given appropriate consent for:

1. Analysis of DNA for diagnostic purposes.
2. Indefinite storage of DNA.
3. Use of anonymous DNA as control samples.

A copy of our consent form is available upon request.