



# Thursdays Webinars



## Recommendations on PKD diagnosis

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Pathophysiology of Anemias Unit

ERN-EuroBloodNet subnetwork Red Blood Cells Disorders

Milan – Italy

8<sup>th</sup> October 2020



Co-funded by  
the Health Programme  
of the European Union



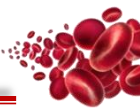
Fondazione IRCCS Ca' Granda  
Ospedale Maggiore Policlinico

Sistema Socio Sanitario



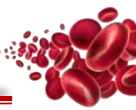
European  
Reference  
Network  
for rare or low prevalence  
complex diseases

Network  
Hematological  
Diseases (ERN EuroBloodNet)



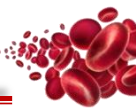
## Conflicts of interest

Consultant at Agios Pharmaceuticals



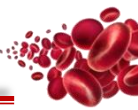
## Practical issues before starting

- ✓ **30-35min presentation (30 slides max) + 15 min Q&A session**
- ✓ **Microphones will be muted by host to avoid back noise**
- ✓ **Please, stop your video to improve internet connexion**
- ✓ **Send your questions during the presentation through the chat, they will be gathered and answered after the presentations.**



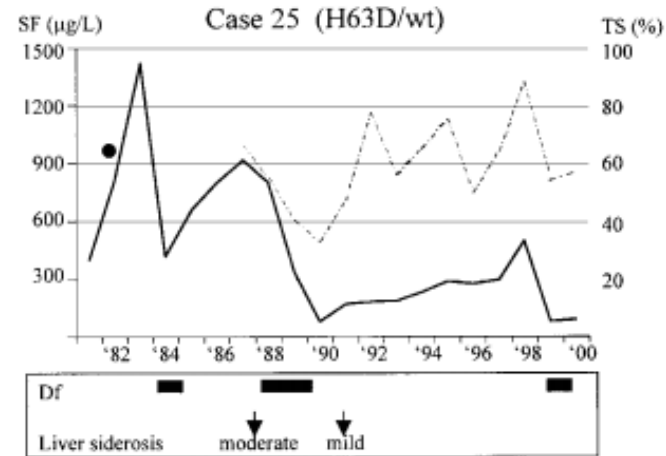
## Learning objectives of the webinar

- ✓ Pathophysiology and phenotypic variability of PK deficiency
- ✓ Diagnostic approaches to PK deficiency
- ✓ The genotype-phenotype correlation in PK deficiency



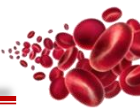
## Case 1

Man	75 yrs
Diagnosis	31 yrs (1975)
Hb	12.5 g/dL
Retic	123 $10^9/L$
Serum ferritin	810 $\mu L$
Splenectomy	No
Transfusions	No
HFE genotype	H63D/wt



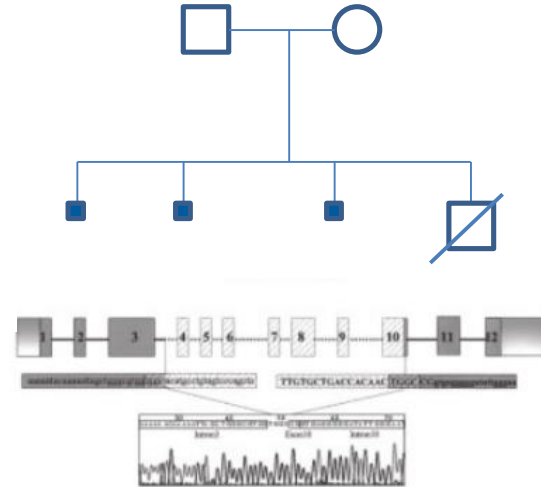
**PKLR genotype** c.1675T/c.1456T

Zanella et al, 2001



## Case 2

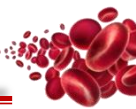
Male 1 day  
Diagnosis at birth  
Death at birth  
Hb 8,9 g/dL  
Retic na  
Serum ferritin >4000 ul/L  
ExTx Yes



## **PKLR** genotype

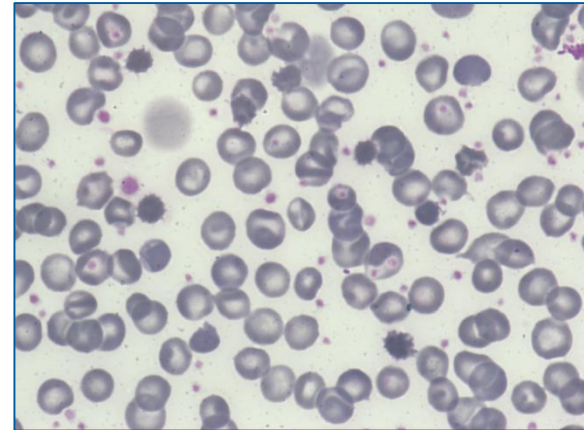
c.409G>A/c.283+1914\_1434del

Fermo et al, 2005



## Case 3

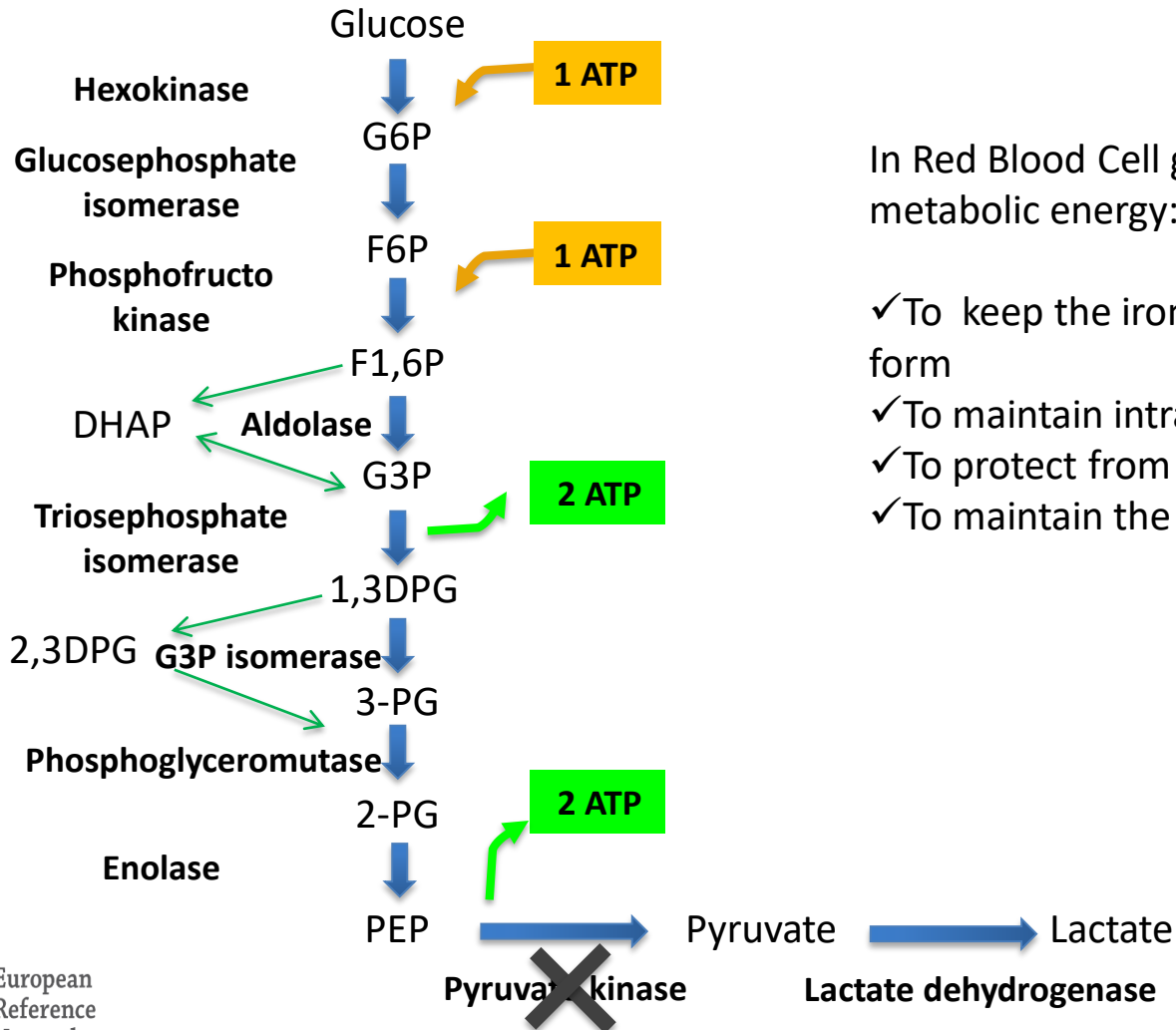
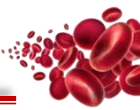
Male,	17 yrs
Diagnosis	13 yrs
Previous diagnosis	CDAll (SEC23B normal)
Hb (birth)	8 g/dL
Hb (1 month)	3g/dL
TX	2U/mo
Splenectomy	8 yrs
Hb (post splen)	8.1g/L
TX (12 yrs)	2U/mo



<b>PK activity</b>	6.0 IU/gHb (11.9-16.7)
<b>PKLR genotype</b>	c.1528C>T/ c.1528C>T

Fermo et al, 2005

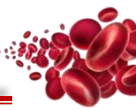
# The Embden-Meyerhof Pathway



In Red Blood Cell glycolysis is the main source of metabolic energy:

- ✓ To keep the iron of hemoglobin in the functional form
- ✓ To maintain intracellular ions concentration
- ✓ To protect from oxydative stress
- ✓ To maintain the red cell shape





Mutation in *PKLR* gene

↓ PK activity  
↑ PK instability



Inefficient glycolysis

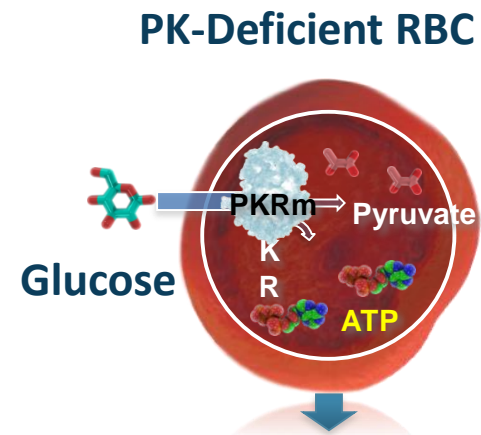
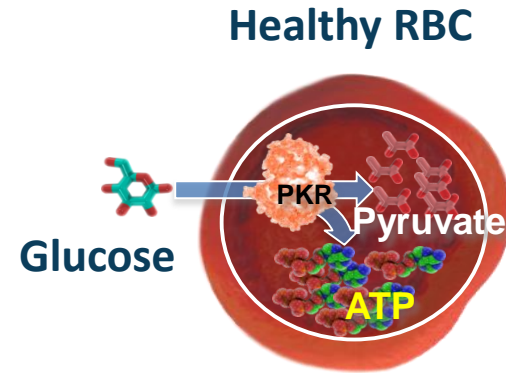
↓ ATP generation



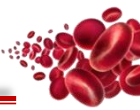
RBC membrane integrity /deformability  
Premature removal from the circulation



Extravasascular hemolysis  
**Chronic hemolytic anemia**



**HEMOLYSIS**



Gene	Isoform	Expression
<b>PKLR</b> CAAT      TATA PKLR Location: 1q21 	<b>PKR (R for Red cell)</b> PKR mRNA: E1 E2 E3E4E5 E12	<ul style="list-style-type: none"> <li>Exclusively in RBCs</li> </ul>
	<b>PKL (L for Liver)</b> PKL mRNA: E2 E3E4E5 E12	<ul style="list-style-type: none"> <li>Liver</li> <li>Kidney</li> <li>Small intestine</li> </ul>
<b>PKM (M for Muscle)</b> PKM Location: 15q22 	<b>PKM1</b> PKM1 mRNA: E1-E7 E8 E9 E11 E12	<ul style="list-style-type: none"> <li>Muscle</li> <li>Brain</li> <li>Heart</li> </ul>
	<b>PKM2</b> PKM2 mRNA: E1-E7 E8 E10 E11 E12	<ul style="list-style-type: none"> <li>Most adult tissues</li> </ul>

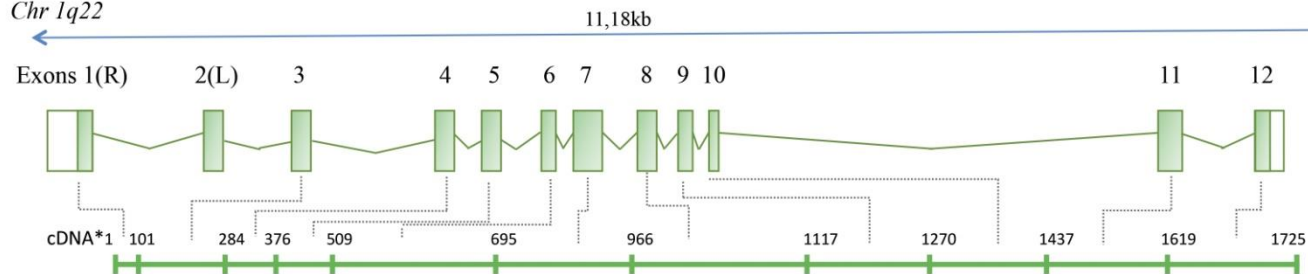
Derived from Israelsen and Semin (2015).

- The *PKLR* gene is located on chromosome 1(1q21)
- Transcription of *PK* tissue-specific promoters yields PKR and PKL
- The cDNA of PKR is 2060 bp long and codes for 574 amino acids
- PKR is expressed exclusively in red blood cells



**A**

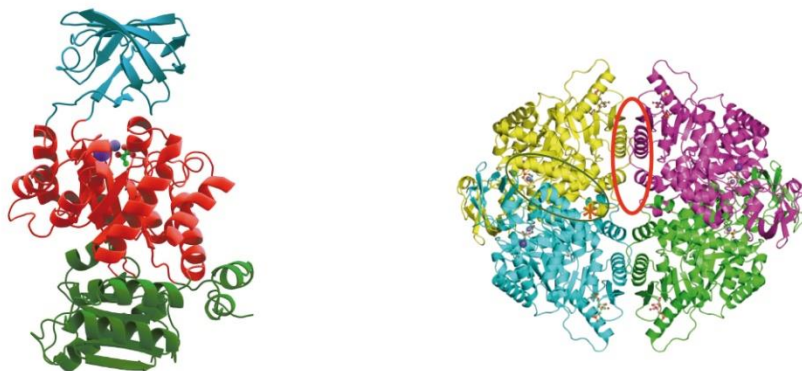
*PKLR gene*  
Chr 1q22

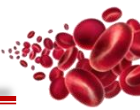


**B**



**C**

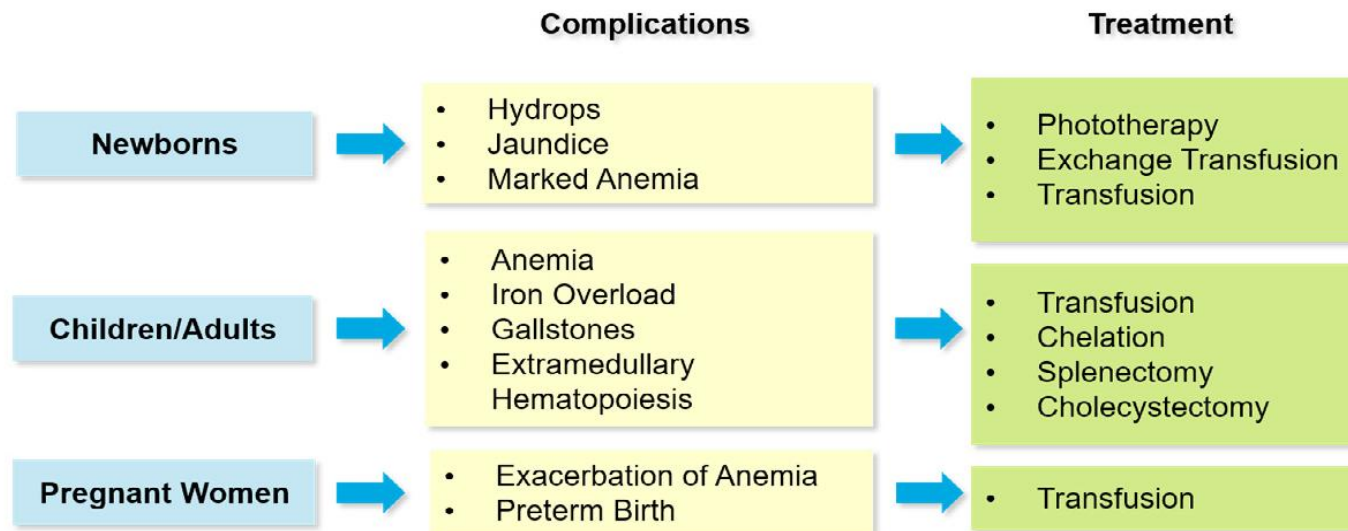




# PK deficiency: a rare form of congenital hemolytic anemia

- ✓ The most common glycolytic defect causing non-spherocytic hemolytic anemia
- ✓ Autosomal recessive inheritance pattern
- ✓ World-wide distribution
- ✓ Heterogeneous clinical presentation

## Clinical presentation and complications



# How rare is PK deficiency?



BLOOD, 1 JUNE 2000 • VOLUME 95, NUMBER 11

RED CELLS

## Estimating the prevalence of pyruvate kinase deficiency from the gene frequency in the general white population

Ernest Beutler and Terri Gelbart

PKD genetic prevalence:  
1:20.000

**Human  
Heredit**

Hum Hered 1992;42:179-183

Original Paper

## Hereditary Nonspherocytic Hemolytic Anemia Due to Pyruvate Kinase Deficiency: A Prevalence Study in Quebec (Canada)

de Medicis E.<sup>a</sup> · Ross P.<sup>a</sup> · Friedman R.<sup>b</sup> · Hume H.<sup>c</sup> · Marceau D.<sup>d</sup> · Milot M.<sup>e</sup> · Lyonnaïs J.<sup>f</sup> · de Braekeleer M.<sup>g</sup>

PKD prevalence in different  
provinces of Quebec:  
1:51,000-1:454,000

To the editor:

BLOOD, 1 DECEMBER 2000 • VOLUME 96, NUMBER 12

## Prevalence of pyruvate kinase deficiency in a northern European population in the north of England

Peter J. Carey, John Chandler, Alex Hendrick, Michael M. Reid, Peter William G. Saunders, Hazel Tinigate, Penelope R. Taylor, and Nicholas West, on behalf of the Northern Region Haematologists Group

PKD prevalence in Northern EU  
populations:  
1:300,000



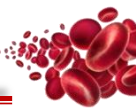
European  
Reference  
Network

for rare or low prevalence  
complex diseases



Network  
Hematological  
Diseases (ERN EuroBloodNet)

Beutler & Gelbart, 2000; De Medicis et al, 1999; Carery et al 2000

Thursdays Webinars

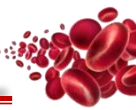


## Prevalence of pyruvate kinase deficiency: A systematic literature review

Matthew H. Secret<sup>1</sup>  | Mike Storm<sup>2</sup> | Courtney Carrington<sup>1</sup> | Deborah Casso<sup>3</sup>  | Keely Gilroy<sup>2</sup> | Leanne Pladson<sup>2</sup> | Audra N. Boscoe<sup>2</sup>

The prevalence of clinically diagnosed PK deficiency is likely between 3.2 and 8.5 per million in Western populations, while the prevalence of diagnosed and undiagnosed PK deficiency could possibly be as high as 51 per million.

**~5-15-fold difference between genetic estimates and diagnosed cases**



✓ **Recessive transmission**

✓ **Lack of knowledge of the disease:**

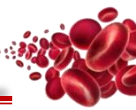
“The diagnosis of PK deficiency is made only when the physician considers the possibility and has red cell enzyme assays performed.” (E. Beutler)

✓ **Heterogeneous clinical phenotype (mild forms - intrauterin death?)**

✓ **Technical problems:**

- Recent transfusions
- WBCs/platelet contamination
- Increased reticulocyte number
- Variants displaying *in vitro* normal enzyme activity

# Is PK deficiency misdiagnosed / underdiagnosed?



[Am J Hematol](#). 2018 May;93(5):672-682. doi: 10.1002/ajh.25058. Epub 2018 Feb 24.

## Multi-gene panel testing improves diagnosis and management of patients with hereditary anemias.

[Russo R](#)<sup>1,2</sup>, [Andolfo I](#)<sup>1,2</sup>, [Manna F](#)<sup>1,2</sup>, [Gambale A](#)<sup>1,2</sup>, [Marra R](#)<sup>1,2</sup>, [Rosato BE](#)<sup>1,2</sup>, [Caforio P](#)<sup>1,2</sup>, [Pinto V](#)<sup>3</sup>, [Pignataro P](#)<sup>2</sup>, [Radhakrishnan K](#)<sup>4,5</sup>, [Unal S](#)<sup>6</sup>, [Tomaiuolo G](#)<sup>7</sup>, [Forni GL](#)<sup>3</sup>, [Iolascon A](#)<sup>1,2</sup>.

“45.5% of the probands originally classified as CDA exhibited a conclusive diagnosis of chronic anemia due to enzymatic defects, mainly due to mutations in *PKLR* gene.”

[Eur J Haematol](#). 2018 Sep;101(3):297-304. doi: 10.1111/ejh.13097. Epub 2018 Jun 25.

## Targeted next generation sequencing for the diagnosis of patients with rare congenital anemias.

[Shefer Averbuch N](#)<sup>1,2</sup>, [Steinberg-Shemer O](#)<sup>1,2</sup>, [Dgany O](#)<sup>3</sup>, [Krasnov T](#)<sup>3</sup>, [Noy-Lotan S](#)<sup>3</sup>, [Yacobovich J](#)<sup>1,2</sup>, [Kuperman AA](#)<sup>4,5</sup>, [Kattamis A](#)<sup>6</sup>, [Ben Barak A](#)<sup>7</sup>, [Roth-Jelinek B](#)<sup>8</sup>, [Chubar E](#)<sup>9</sup>, [Shabad E](#)<sup>10</sup>, [Dufort G](#)<sup>11</sup>, [Ellis M](#)<sup>2,12</sup>, [Wolach O](#)<sup>2,13</sup>, [Pazgal I](#)<sup>2,14</sup>, [Abu Quider A](#)<sup>15</sup>, [Miskin H](#)<sup>16,17</sup>, [Tamary H](#)<sup>1,2</sup>.






“Genetic diagnosis was achieved in 13 out of 21 patients (62%). Six patients were diagnosed with pyruvate kinase deficiency (28,5%). The mean lag time from presentation to diagnosis was over 13 years.”





**TEST OF THE MONTH**

**Addressing the diagnostic gaps in pyruvate kinase deficiency:  
Consensus recommendations on the diagnosis of pyruvate  
kinase deficiency**

Paola Bianchi<sup>1</sup>  | Elisa Fermo<sup>1</sup> | Bertil Glader<sup>2</sup> | Hitoshi Kanno<sup>3</sup> | Archana Agarwal<sup>4</sup>  |  
Wilma Barcellini<sup>1</sup>  | Stefan Eber<sup>5</sup> | James D. Hoyer<sup>6</sup> | David J. Kuter<sup>7</sup>  |  
Tabita Magalhães Maia<sup>8</sup> | Maria del Mar Mañu-Pereira<sup>9</sup> | Theodosia A. Kalfa<sup>10</sup> |  
Serge Pissard<sup>11</sup> | José-Carlos Segovia<sup>12,13</sup> | Eduard van Beers<sup>14</sup>  | Patrick G. Gallagher<sup>15</sup> |  
David C. Rees<sup>16</sup> | Richard van Wijk<sup>17</sup> | with the endorsement of EuroBloodNet, the  
European Reference Network in Rare Hematological Diseases

Global PK deficiency International expert group (2016)  
(24 experts from 20 different Expert Centres )



Survey on diagnostic methodologies



Forum discussion

7 Centres from EU, 5 from USA, and 1 from Asia



**Consensus diagnostic recommendations  
Algorithm for the diagnosis of PK deficiency**

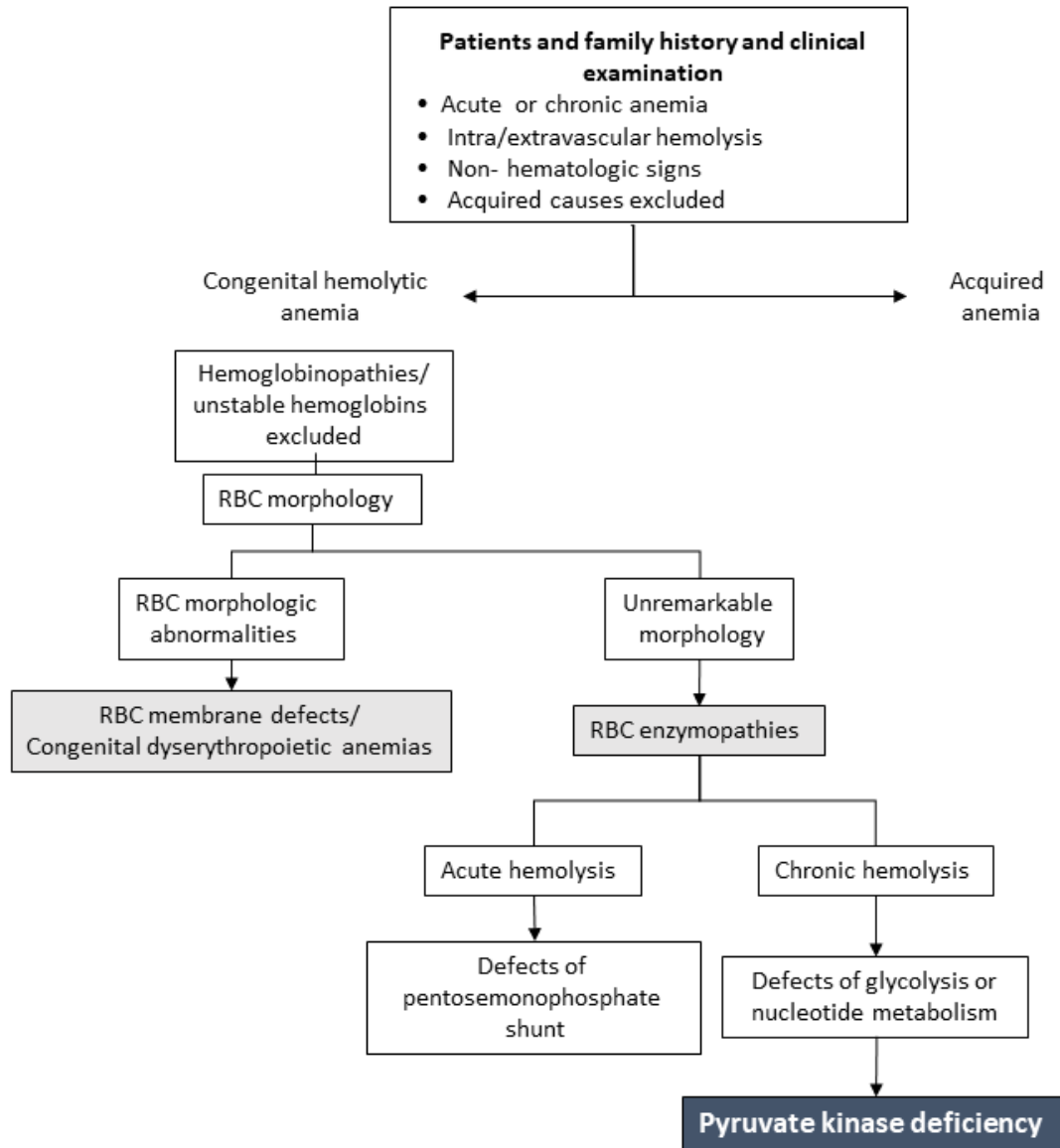
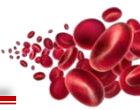


TEST OF THE MONTH

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Paola Bianchi<sup>1</sup> | Elisa Fermo<sup>1</sup> | Bertil Glader<sup>2</sup> | Hitoshi Kanno<sup>3</sup> | Archana Agarwal<sup>4</sup> | Wilma Barcellini<sup>1</sup> | Stefan Eber<sup>5</sup> | James D. Hoyer<sup>6</sup> | David J. Kuter<sup>7</sup> | Tabita Magalhães Maia<sup>8</sup> | Maria del Mar Mañu-Pereira<sup>9</sup> | Theodosia A. Kalfa<sup>10</sup> | Serge Pissard<sup>11</sup> | José-Carlos Segovia<sup>12,13</sup> | Eduard van Beers<sup>14</sup> | Patrick G. Gallagher<sup>15</sup> | David C. Rees<sup>16</sup> | Richard van Wijk<sup>17</sup> | with the endorsement of EuroBloodNet, the European Reference Network in Rare Hematological Diseases

	Recommendation	Evidence
<i>Clinical presentation</i>	PK deficiency may be suspected in: <ul style="list-style-type: none"> <li>- patients with variable chronic anaemia and/or splenomegaly and/or jaundice, with normal or near-normal red cell morphology.</li> <li>- transfusion dependent cases of unknown aetiology</li> <li>- haemolytic patients with unexplained severe neonatal indirect hyperbilirubinemia</li> <li>- presence of high reticulocyte number in splenectomised patients with no diagnosis</li> </ul>	Mean: 95% Median: 100% (75-100)
<i>Clinical data</i>	-Information on clinical history (both recent as well as from infancy, ie neonatal jaundice), family history should always be requested together with samples, as well as the time of last blood transfusion	Mean: 98.6% Median:100% (90-100)
<i>Laboratory data</i> <i>(mandatory in bold)</i>	- <b>Complete blood count</b> - <b>RBC morphology</b> -Markers of haemolysis ( <b>reticulocyte count, LDH, unconjugated bilirubin, haptoglobin<sup>1,2</sup></b> )	Mean: 97% Median:100% (90-100)
<i>Differential diagnosis</i>	Acquired haemolytic anaemia, membranopathies, CDAs, unstable haemoglobins, red cell enzymopathies other than PK deficiency should be excluded (See Figure 5)	Mean: 92.1% Median: 100% (50-100)

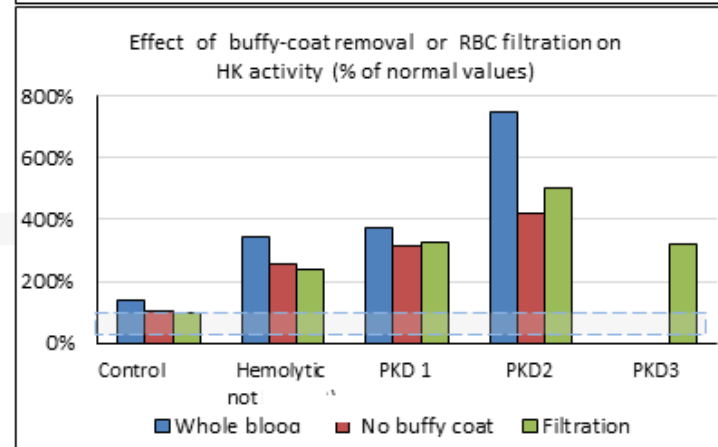
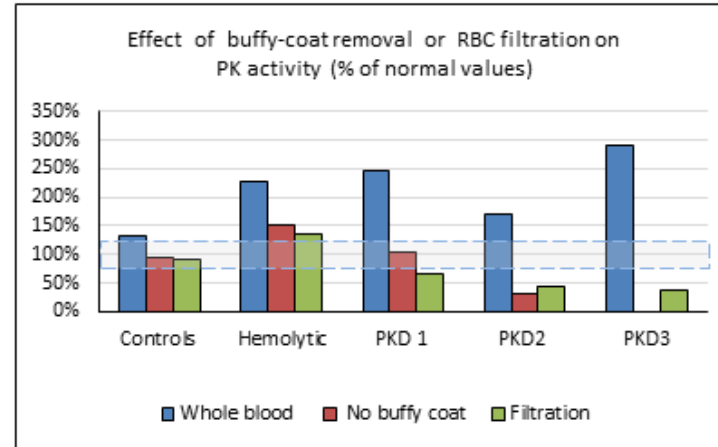
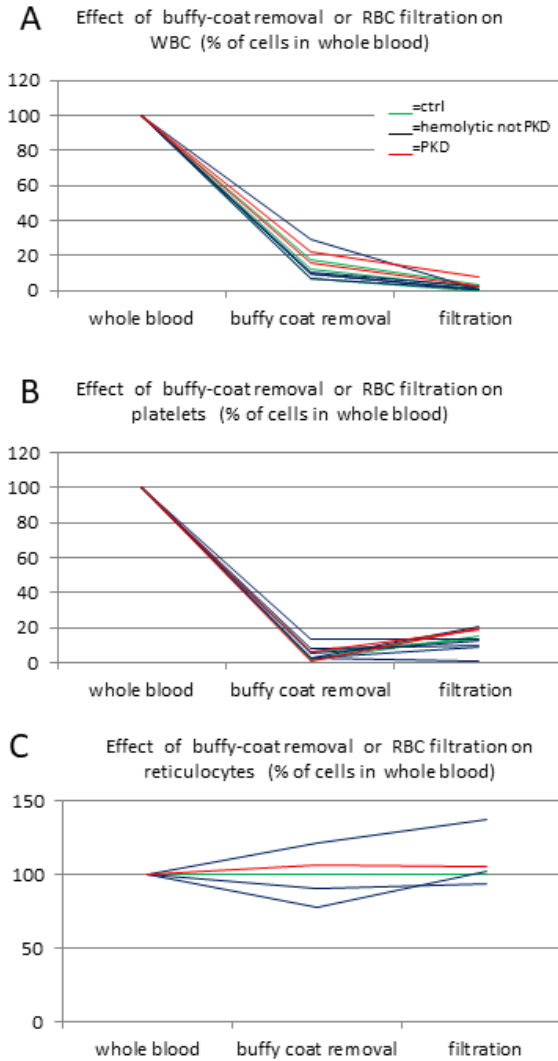
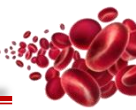


# Laboratory diagnosis of PK deficiency

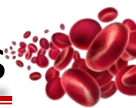


<b>Biochemical testing</b>		
Reference test for biochemical assay	RBC PK activity assay by spectrophotometry (Beutler, 84)	Mean: 98.7% Median: 100% (80-100)
Storage time of sample	PK enzyme assay may be considered stable at 4° C until up to 21 days after collection <sup>3</sup> . A maximum of 14 days storage is recommended if PK activity is related to HK activity due to different stability of HK activity	Mean: 95% Median: 100% (80-100)
Sample anticoagulant	ACD; EDTA, CPD, Heparin could be considered for the enzyme assay (Beutler, 84): EDTA is the main anticoagulant used in daily practice.	Mean: 100% Median: 100%
Sample preparation	Purification on α-cellulose/microcrystalline cellulose column is recommended. Buffy coat removal may be considered as an alternative. PK enzyme activity cannot be performed on whole blood	Mean: 96.7% Median: 100% (80-100)
Reticulocytes interference	Reticulocyte number must be taken into account when interpreting results of PK enzyme assay, particularly when of low-normal PK activity levels. Results could be compared with enzyme activities obtained from a control sample with the same degree of reticulocytosis, or by calculating the ratio of PK activity to another cell age dependent enzyme (e.g. hexokinase).	Mean: 96.1% Median: 100% (70-100)
Interference of donor red blood cells	The enzyme assay should be performed as far as possible after a red cell transfusion. The laboratory should record the time since transfusion. A minimum of 50 days from last transfusion is considered a "safe" period for testing of PK activity, leading to an estimated donor RBC contamination of about 7-14%. Results of enzyme activity need to be interpreted with caution in transfused patients <sup>4</sup> .	Mean: 96.9% Median: 100% (60-100)
Confirmatory tests	In case of decreased PK activity, sequencing of <i>PKLR</i> gene is highly recommended to confirm the diagnosis	Mean: 88.3% Median: 100% (10-100)

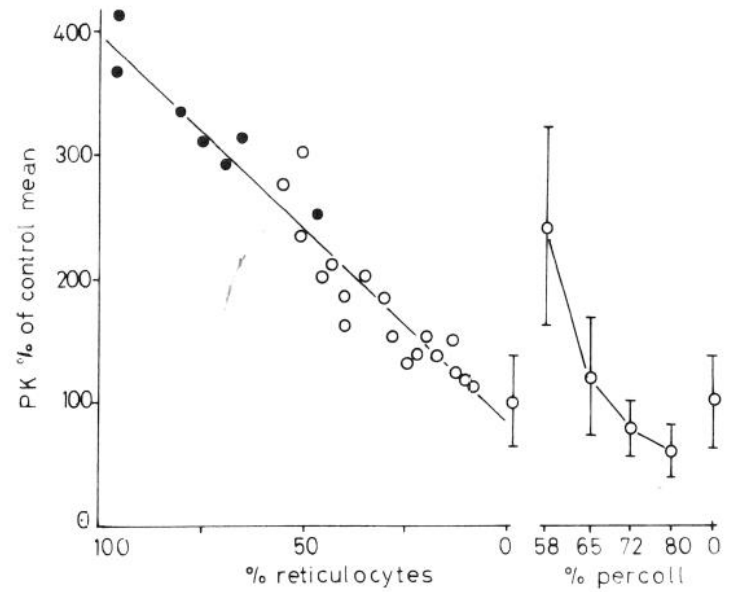
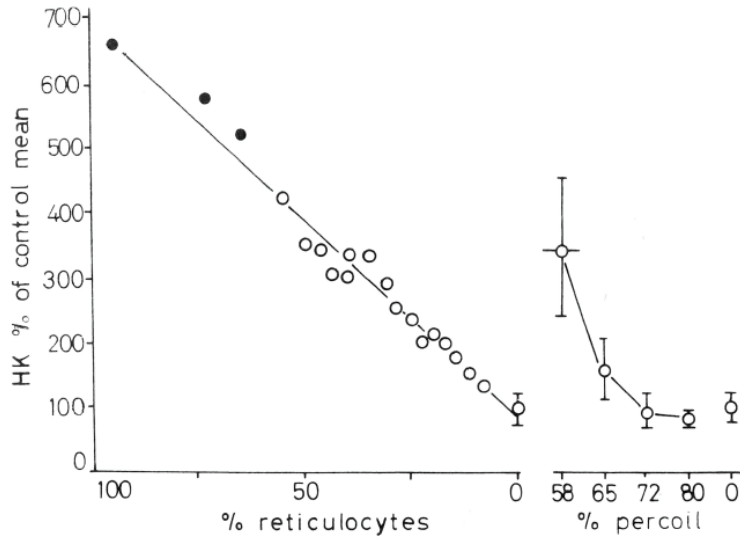
# Laboratory diagnosis of PK deficiency – WBC influence



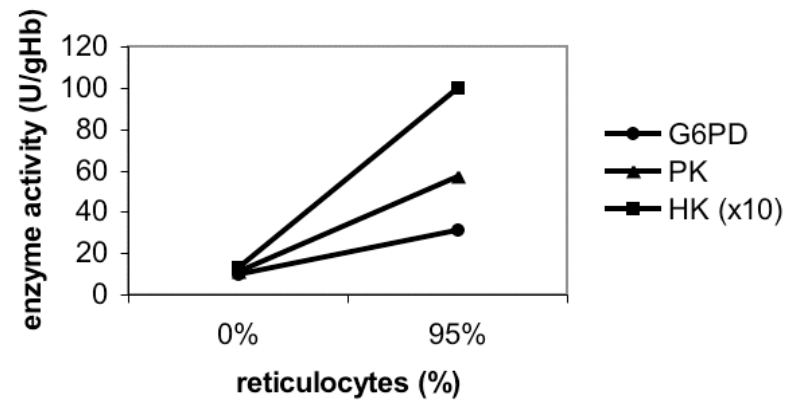
# Laboratory diagnosis of PK deficiency – Influence of reticulocytosis

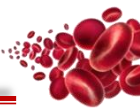


Age Dependence Behaviour of Red Cell Glycolytic Enzymes



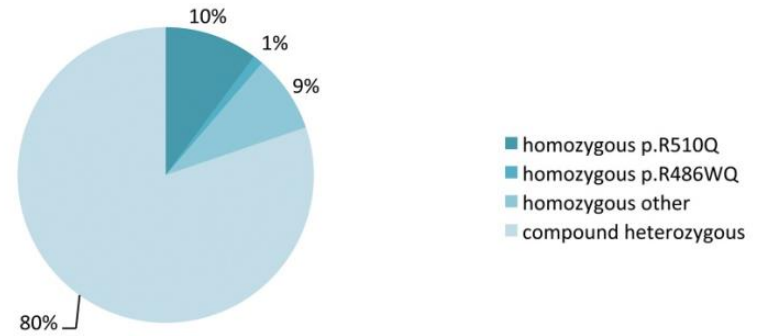
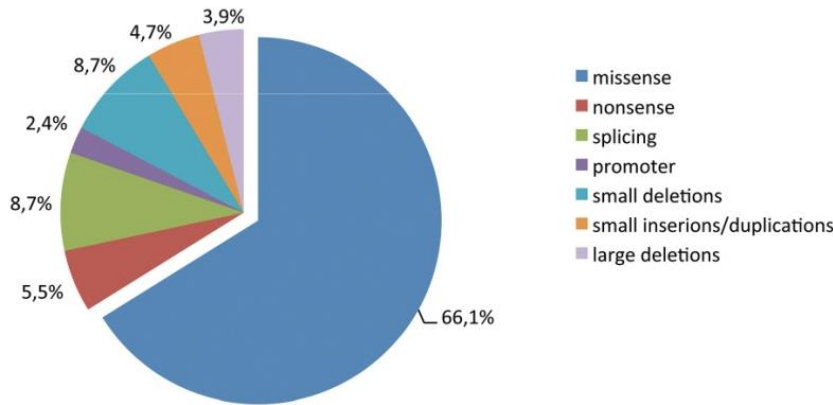
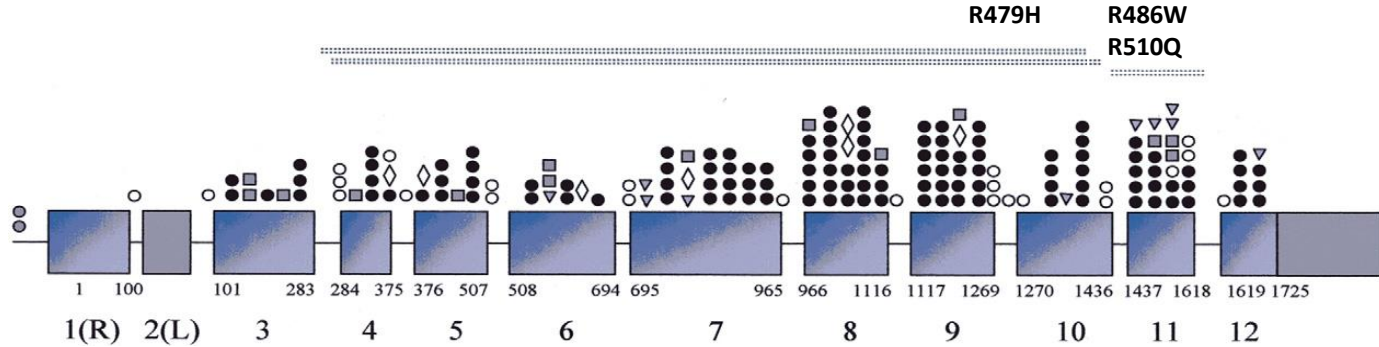
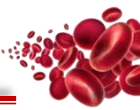
Jansen et al. Br J Haematol 1985; 61: 51-59





<b>Molecular testing</b>		
<i>Indication</i>	<p>-Molecular testing is highly recommended to confirm a suspected case of PK deficiency based on decreased enzyme activity.</p> <p>-Molecular testing of <i>PKLR</i> gene by Sanger is suitable for patients with (relatively) decreased PK activity</p> <p>- Use of NGS panels is a reliable alternative method for diagnosis of PK deficiency. It is particularly relevant for:</p> <ul style="list-style-type: none"> <li>- neonates (if family study is not available)</li> <li>- transfusion dependent patients/recently transfused patients</li> <li>- samples with prolonged shipping times</li> </ul>	<p>Mean: 91.2%</p> <p>Median: 100%</p> <p>(10-100)</p>
<i>PKLR genotype discrepancies</i>	<p>In case of genotype discrepancies (patients with suspected PKD and one or none mutations detected) further investigation are required:</p> <p>-Assays for detection of large deletions</p> <p>-Re-evaluation of other causes of haemolysis by specific tests or NGS platform</p> <p>In absence of any mutation and decreased PK activity:</p> <p>- NGS tools or, <i>KLF1</i> gene mutations should be considered</p>	<p>Mean: 92.5%</p> <p>Median: 100%</p> <p>(40-100)</p>

# Laboratory diagnosis of PK deficiency – Molecular testing



## Unknown intronic variants

Lezon-Geyda K, Rose MJ, McNaull MA, et al. Pklr Intron Splicing-Associated Mutations and Alternate Diagnoses Are Common in Pyruvate Kinase Deficient Patients with Single or No PKLR Coding Mutations. ASH Meeting 2018



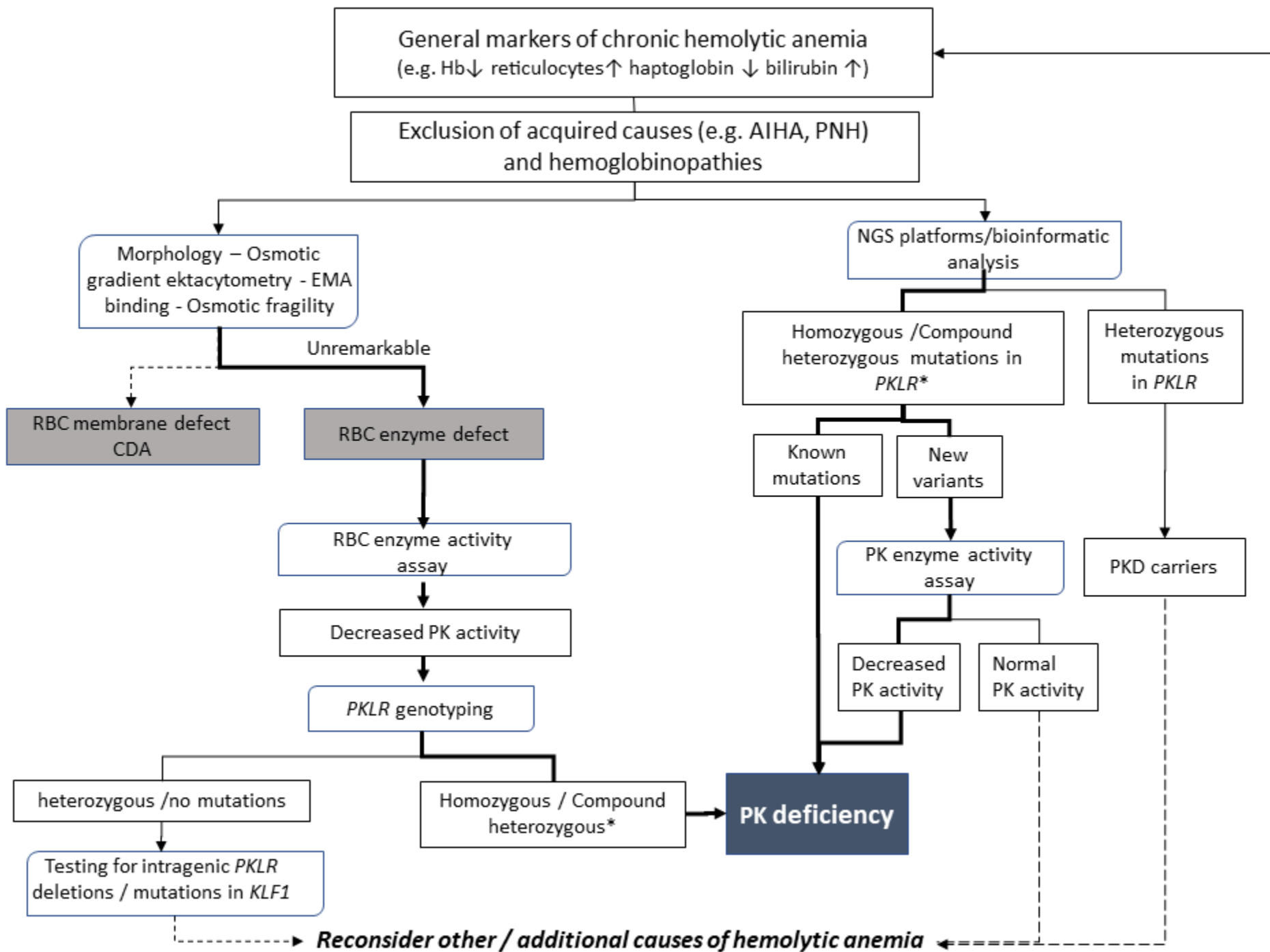
# Laboratory diagnosis of PK deficiency – NGS



Reference	Method	N. of genes analyzed	N. of cases studied with CHA	PKD diagnosis	New diagnosis and number and type of mismatched diagnoses
Svidnicki et al, 2018	t-NGS	35	36	2	2 new PKD
	t-NGS	55	43	8	8 new PKD
Jamwal et al, 2020	WES	n.a.	4	4	4 new PKD
	t-NGS	76	21 <sup>a</sup>	6	3 new PKD
Qin et al, 2020					2 CDA → PKD
					1 DBA → PKD
Kedar et al, 2019	t-NGS	76	21 <sup>b</sup>	6	4 new PKD
					2 CDA → PKD
	t-NGS	34 and 71	74 <sup>c</sup>	7	7 CDA → PKD
Shefer Averbuch, 2018					2 new PKD
					1 CDA → PKD

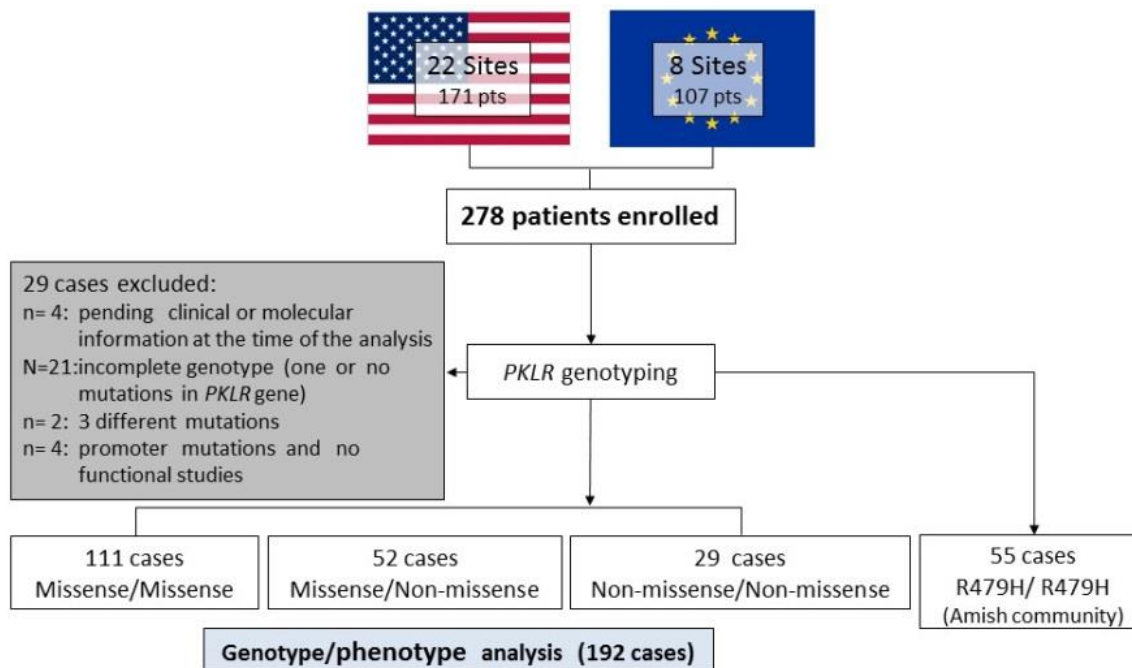
Number of genes included in the panel, number of cases analyzed in each study and cases diagnosed with pyruvate kinase deficiency are shown. Next-generation sequencing analysis allowed modification of a previous diagnosis; the number and the type of mismatched diagnosis is reported in the last column. <sup>a</sup>All transfusion-dependent patients. <sup>b</sup>No diagnosis despite extensive laboratory investigations. <sup>c</sup>Suspected diagnosis of congenital dyserythropoietic anemia. CHA: chronic hemolytic anemias; PKD: pyruvate kinase deficiency; t-NGS: targeted next-generation sequencing; WES: whole-exome sequencing; n.a.: not available; CDA: congenital dyserythropoietic anemia; DBA: Diamond-Blackfan anemia.

Russo R, et al  
2018  
Roy et al,  
2016

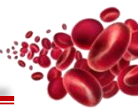


\* *In trans* nature of mutations to be confirmed by family studies

# Genotype-Phenotype Correlation and Molecular Heterogeneity



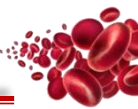
# Genotype-Phenotype Correlation



	NM/NM, N=29 Median (Range) n=29	M/NM, N=52 Median (Range) n=50	M/M, N=111 Median (Range) n=105	p-value <sup>+</sup>
Age at diagnosis (years)	0.4 (0-10.9) n=29	0.7 (0-42.3) n=50	1.3 (0-60.3) n=105	0.049
Hemoglobin (g/dl)**	7.9 (6.5-8.9) n=14	8.4 (6.4-12.8) n=21	9.2 (4.3-12.3) n=40	0.003*
Total number of lifetime transfusions	65 (3-991) n=27	25 (1-721) n=38	16 (1-1915) n=81	0.0013*
Maximum ferritin (ng/ml)	1787 (423-13,409) n=22	604 (22-8,220) n=37	573 (31-9,679) n=75	<0.0001*
PK enzyme activity normalized to patient-specific normal range (%)	-41.6 (-152.4-15.2) n=18	-51.9 (-211.1-64.4) n=24	-69.6 (-485.7-117.6) n=60	0.16

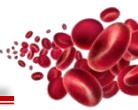
## Response to splenectomy

		No response (Hb <8 g/dl) n=31	Partial response (Hb 8-<11 g/dl) n=110	Complete response (Hb ≥11 g/dl) n=7	p
<b>Genotype</b>	<b>M/M</b>	29%	59%	100%	0.0017
	<b>M/NM</b>	32%	26%	0%	0.0005
	<b>NM/NM</b>	39%	16%	0%	0.5

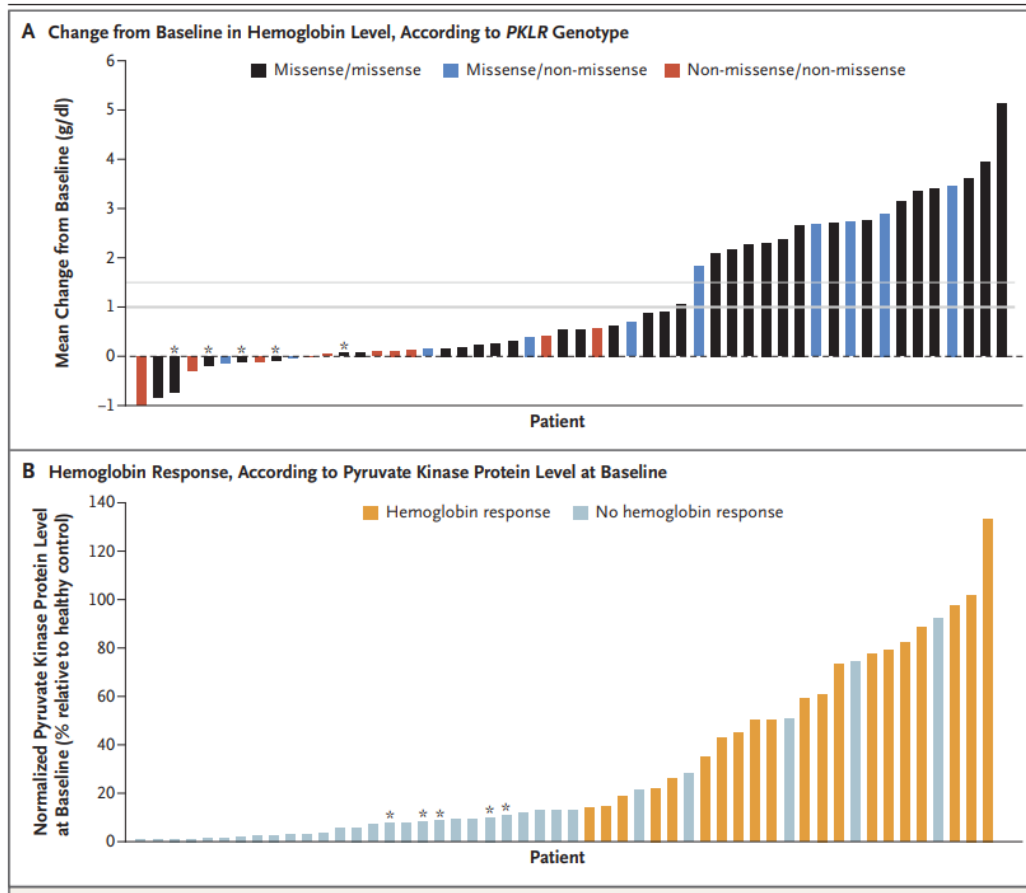


## Perinatal Course and Pregnancy Outcomes of Patients by *PKLR* Mutation Type

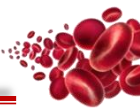
	Missense/Missense N=113 patients	Missense/ Non-Missense N=52 patients	Non-Missense/ Non-Missense N=30 patient	P value**
<b>Characteristics</b>	<b>n</b>	<b>n</b>	<b>n</b>	
<b>Perinatal</b>	28/100 (28%)	15 /48 (31%)	11/28 (39%)	0.26
<b>Complications/Treatment</b>				
<b>In utero transfusions</b>	12/26 (46%)	8/15 (53%)	4/10 (40%)	1
<b>Hydrops</b>	3/26 (12%)	5/14 (36%)	1/11 (9%)	1
<b>Exchange transfusion</b>	34/80 (42%)	17/40 (43%)	9/21 (43%)	1
<b>Pregnancy Outcomes/Management (data presented per pregnancy)*</b>	n=29 pregnancies* 16 female patients	n=13 pregnancies* 6 female patients	n=7 pregnancies* 3 female patients	
<b>Normal birth - Full-term</b>	20/29 (69%)	8/13 (62%)	5/7 (71%)	1
<b>Pre-maturity</b>	3/29 (10%)	0/13 (0%)	2/7 (29%)	0.24
<b>Transfusions during pregnancy</b>	9/21 (43%)	3/6 (50%)	5/5 (100%)	0.043



## Safety and Efficacy of Mitapivat in Pyruvate Kinase Deficiency



# Genotype-phenotype correlation in patients underwent BMT



	Sex	Country	Genotype	Mutation effect		Splenectomy	Age at HCST	Year	Outcome	Structure ref
Pt 1	M	Asia	Unknown	Unknown	Unknown	No	5 y	1996	Alive	
Pt 2	F	EU	p. [E241*; R532W]	Nonsense	<b>Missense</b> <sup>1</sup>	Yes	15	2002	Deceased	10,11
Pt 3	F	Asia	p.[K348N; R359H]	<b>Missense</b> <sup>2</sup>	<b>Missense</b> <sup>2</sup>	No	1 y 7 mo	2009	Alive	47
Pt 4	F	EU	p.[E241*; R488Q]	Nonsense	Missense	No	3 y	2009	Alive	
Pt 5	M	Asia	p. [R40Q; D339N]	Missense	<b>Missense</b> <sup>3</sup>	No	2 y 6 mo	2009	Alive	10
Pt 6	F	EU	p. [M377fs; M377fs]	Nonsense	Nonsense	Yes	17 y	2010	Deceased	
Pt 7	F	EU	p.[G165V; R510Q]	Missense	<b>Missense</b> <sup>4</sup>	Yes	39 y	2011	Deceased	16
Pt 8	F	EU	p.[G511E; E538*]	Missense	Nonsense	Yes	7 y	2013	Alive	
Pt 9	M	EU	p.[I494T; R559*]	Missense	Nonsense	No	6 y	2013	Deceased	
Pt 10	M	Asia	p.[V283A; I314T]	Missense	<b>Missense</b> <sup>3</sup>	No	1 y 6 mo	2013	Alive	10
Pt 11	M	EU	p.[K541fs; K541fs]	Nonsense	Nonsense	Yes	10 y	2014	Deceased	
Pt 12	M	Asia	p.[D221Y; I314T]	<b>Missense</b> <sup>3</sup>	<b>Missense</b> <sup>3</sup>	No	9 y	2014	Alive	10,17
Pt 13	M	Asia	p.[V283A; V283A]	Missense	Missense	No	1 y 6 mo	2015	Alive	
Pt 14	M	EU	p.[D331Q; D339H]	<b>Missense</b> <sup>5</sup>	<b>Missense</b> <sup>3</sup>	Yes	41 y	2015	Alive	10,14
Pt 15	M	Asia	c.[1270-3C>A]; p.[G540*]	Nonsense	Nonsense	Yes	11 y	Unknown	Alive	
Pt 16	F	Asia	c.[1270-3C>A]; p.[G540*]	Nonsense	Nonsense	No	8 y	Unknown	Alive	
Ref 94	F	China	p.[I314T; I314T]	<b>Missense</b> <sup>3</sup>	<b>Missense</b> <sup>3</sup>	No	Unknown	Unknown	Alive	
Ref 95	M	Japan	p.[Pro145Hisfs; Pro145Hisfs]	Nonsense	Nonsense	Yes	32 y	Unknown	Alive	

Missense variants falling in "strategic" functional amino acid residues or associated with documented thermo-unstable variants are reported in bold. <sup>1</sup>Directly involved in the fructose 1,6 biphosphate activator. <sup>2</sup>Directly involved in the substrate and cation binding sites. <sup>3</sup>Residues directly involved in the allosteric site and catalytic center. <sup>4</sup>Highly unstable. <sup>5</sup>Proximity of the substrate-binding site.

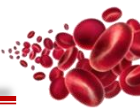
## New PK scheme proposal: UK NEQAS

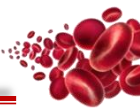
- European collaboration: essential because of small numbers of laboratories in each country
- Performance assessment for quantitative assay
- Could develop to include molecular methods
- Development phases:
  - Survey material development
    - Storage, stability, volumes etc.
  - Recruitment of interested participants
  - Small scale survey with selected labs
  - Pilot exercise(s) to refine scheme design
  - Development of performance assessment methods

**UK NEQAS**  
International Quality Expertise









## Take home messages

1. PKD diagnosis: to monitor complications; to have access to new therapies
2. Biochemical testing and molecular characterization are complementary approaches, not alternative
3. Always consider differential diagnosis
4. Careful costs/benefits evaluation before choosing the diagnostic approach

### Biochemical assay:

😊 € / 2 hours

😞 Other laboratory tests  
to exclude other conditions



### NGS:

😞 €€-€€€ / 15-30 days

😊 Diagnosis of wide spectrum of defects