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Article type : Short Reports

The Pyruvate Kinase (PK) to Hexokinase Enzyme Activity Ratio and PK-R Protein Level in the Diagnosis and Phenotype of PK Deficiency

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This is the author manuscript accepted for publication and has undergone full peer review but has not been through the copyediting, typesetting, pagination and proofreading process, which may lead to differences between this version and the [Version of Record](#). Please cite this article as [doi: 10.1111/BJH.16724](https://doi.org/10.1111/BJH.16724)

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22 **Sources of support (financial, equipment, drugs):** This study was supported by funding from
23 Agios Pharmaceuticals.

24 **Running title:** PK:HK Ratio in Diagnosis and PK-R in PKD Phenotype

25 **Figures:** 1

26 **Tables:** 1

27 **References:** 10

28 **Key words:** Pyruvate kinase, PKD, hexokinase, diagnosis, hemolysis, enzyme assay, pyruvate
29 kinase protein, congenital hemolytic anemia

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31

32 **ABSTRACT**

33

1 Diagnosis of pyruvate kinase deficiency (PKD), the most common cause of
2 hereditary nonspherocytic hemolytic anemia, remains challenging in routine practice
3 and no biomarkers for clinical severity have been characterized. This prospective study
4 enrolled 41 patients with molecularly-confirmed PKD from 9 North American centers to
5 evaluate the diagnostic sensitivity of pyruvate kinase (PK) enzyme activity and
6 PK:hexokinase (HK) enzyme activity ratio and evaluate the pyruvate kinase-red cell
7 (PK-R) protein level and erythrocyte metabolites as biomarkers for clinical severity. In
8 this population not transfused for ≥ 90 days before sampling, the diagnostic sensitivity of
9 PK enzyme assay was 90% (95% CI, 77%-97%) whereas the PK:HK ratio sensitivity
10 was 98% (95% CI, 87%-100%). There was no correlation between PK enzyme activity
11 and clinical severity. Transfusion requirements correlated with normalized erythrocyte
12 ATP levels ($r=0.527$, $P=0.0016$) and PK-R protein levels ($r=-0.527$, $P=0.0028$). PK-R
13 protein levels were significantly higher in never transfused (median=40.1%, range=9.8-
14 73.9%) versus ever transfused (median=7.7%, range=0.4-15.1%) patients ($P=0.0014$).
15 The PK:HK ratio had excellent sensitivity for PK diagnosis, superior to *PKLR* exon
16 sequencing. Given that the number of *PKLR* variants and genotype combinations limits
17 prognostication based on molecular findings, PK-R protein level may be a useful
18 prognostic biomarker of disease severity and merits further study.

19 **INTRODUCTION**

20 Pyruvate kinase deficiency (PKD) is the most common cause of chronic
21 hereditary non-spherocytic hemolytic anemia and results from mutations in the *PKLR*
22 gene. Over 300 distinct pathogenic mutations have been documented (Pissard, *et al*
23 2006, Warang, *et al* 2013). Prevalence estimates of PKD vary widely between 1:20,000
24 and 1:300,000 in Caucasian populations (Beutler and Gelbart 2000, Carey, *et al* 2000),
25 likely because the diagnosis of PKD remains challenging in routine practice (Al-
26 Samkari, *et al* 2020). While the pyruvate kinase (PK) enzyme assay is typically
27 employed as an initial diagnostic screen in patients with unidentified Coombs-negative
28 hemolytic anemia, PK enzyme levels may be falsely normal in patients receiving red cell
29 transfusion within 90 days, in those with profound reticulocytosis (as PK enzyme activity
30 is red cell age-dependent), or in patients with *PKLR* mutations resulting in unusual
31 biochemical properties (Bianchi, *et al* 2019, Grace, *et al* 2015). Molecular testing

1 similarly presents challenges: up to 20% of affected patients have new *PKLR* variants,
2 up to 10% of patients have variants not detected through routine exon sequencing, and
3 molecular testing may be cost-prohibitive (Grace, *et al* 2018). After the diagnosis is
4 established, there are no known biomarkers of clinical severity which could be utilized
5 for determining prognosis or monitoring. While PK enzyme activity does not have a
6 relation with *PKLR* mutation severity (Grace, *et al* 2018), it remains unresolved whether
7 phenotypic severity has any relation to enzyme activity in PKD. Given the current
8 diagnostic limitations and the broad phenotypic spectrum of disease, alternative
9 diagnostic strategies and biomarkers of clinical severity are needed.

10 Therefore, we aimed to describe the correlation of PK enzyme activity,
11 erythrocyte PK (PK-R) protein level, and erythrocyte metabolites [adenosine
12 triphosphate (ATP), 2,3-diphosphoglycerate (2,3-DPG)] with clinical phenotype as well
13 as estimate the sensitivity of PK enzyme activity and the pyruvate kinase:hexokinase
14 (PK:HK) ratio to diagnose PKD in a cohort of patients with genetically-confirmed PKD.
15 Because hexokinase (HK) is also red cell age-dependent, the use of the PK:HK ratio
16 may correctly diagnose patients with normal PK enzyme activity in the setting of
17 profound reticulocytosis, a common finding in PKD. This ratio has previously been
18 demonstrated to correctly diagnose PK deficiency in a few select patients (Bianchi, *et al*
19 2019). In this study, we had the opportunity to establish the validity of this observation in
20 a large number of patients with molecularly-confirmed PKD.

21 22 **METHODS**

23 A subset of 9 North American sites from the PKD Natural History Study (NHS)
24 (NCT02053480) (Grace, *et al* 2018) participated in a sub-study, in which blood samples
25 were collected at a single time-point from all enrolled patients. All patients had two
26 confirmed *PKLR* mutations and no red cell transfusions for ≥ 90 days prior to the blood
27 sample. Erythrocyte PK and HK enzyme activity testing were performed using standard
28 biochemical assays (Beutler 1984) at the Stanford Red Blood Cell Special Studies
29 Laboratory (reference range for PK activity, 3.2-6.5 EU/gm Hb and for HK activity, 0.14-
30 0.37 EU/gm Hb). Baseline erythrocyte PK-R protein was quantitated by antibody-based
31 capture and detection using an electrochemiluminescence immunoassay (Meso Scale

1 Discovery, Rockville, MD) and the signal normalized to a control sample without PKD.
2 Erythrocyte ATP and 2,3-DPG concentrations ($\mu\text{g/ml}$) in blood were analyzed using
3 qualified tandem mass spectrometry methods and then normalized for individual
4 hemoglobin (Hb) values [ATP and 2,3-DPG concentrations were converted to g/dL and
5 divided by Hb (g/dL)]. Spearman correlation coefficients were calculated to estimate the
6 correlation between continuous variables, and the Wilcoxon rank-sum test was used to
7 assess the association of continuous and binary variables.

8

9 **RESULTS AND DISCUSSION**

10 Of the 255 patients in the PKD NHS, 41 patients participated in this sub-study.
11 The median age was 25.3 years (range 1.4-60.4) and 80% of patients were
12 splenectomized (Table). Mean measured PK enzyme activity level was 1.1 EU/gm Hb;
13 37/41 of patients had low PK activity, corresponding to a diagnostic sensitivity of 90%
14 [95% confidence interval (CI), 77%-97%]. Normal PK activity was observed in 4
15 patients, 3 of whom had high HK activity and a low PK/HK ratio (reference normal mean
16 PK/HK ratio 15.6, range: 8.7-22.5). In total, 40/41 patients had a low PK/HK ratio,
17 corresponding to a sensitivity for PKD diagnosis of 98% (95% CI, 87%-100%). The
18 single patient not accurately diagnosed with this method had a PK enzyme activity of
19 4.6 EU/gm Hb and a PK/HK ratio of 9.2, which is at the low end of the normal range.
20 There were no correlations between PK enzyme activity and clinical severity indicators,
21 including post-splenectomy hemoglobin ($r=0.109$, $P=0.56$), total number of lifetime
22 transfusions ($r=-0.107$, $P=0.54$), transfusion status (ever transfused vs. never
23 transfused; $P=0.30$), or splenectomy status (splenectomized vs. not splenectomized;
24 $P=0.41$).

25 Normalized ATP levels strongly positively correlated with normalized 2,3-DPG
26 levels ($r=0.93$, $P<0.0001$, Figure 1A), moderately positively correlated with total lifetime
27 transfusions ($r=0.527$, $P=0.0016$, Figure 1B) and percent reticulocyte count ($r=0.581$,
28 $P=0.0003$, Figure 1C), and were higher in ever transfused (median=0.0022,
29 range=0.0011-0.0029) compared with never transfused patients (median=0.0012,
30 range=0.0008-0.0020), ($P=0.0037$).

1 PK-R protein percentage was moderately inversely correlated with the total
2 number of transfusions prior to enrollment ($r=-0.527$, $P=0.0028$, Figure 1D) and was
3 higher in never transfused (median=40.1%, range=9.8-73.9%) compared with ever
4 transfused patients (median=7.7%, range=0.4%-15.1%), ($P=0.0014$, Figure 1E). The
5 PK-R protein level in the Amish population, which shares a common homozygous
6 R479H genotype, was relatively consistent among patients as compared to the wide
7 variation of PK-R protein levels of non-Amish patients with differing genotypes (Table).
8 In addition, PK-R protein levels were numerically higher in patients with two missense
9 *PKLR* variants (median=7.7%) as compared with those with at least one drastic variant
10 (median=2.0%), although this difference was not statistically significant ($p=0.36$).

11 There are two important conclusions from this study. The first is that the PK:HK
12 ratio appears to have excellent sensitivity for the diagnosis of PKD. With a sensitivity of
13 98% in this study population of non-transfused patients, this ratio was more sensitive
14 than PK activity in isolation and even more sensitive than *PKLR* exon sequencing to
15 diagnose PKD (Grace, *et al* 2018). Therefore, based on this data, the PK:HK ratio may
16 be considered a standard component of the PKD diagnostic evaluation. Individuals who
17 are heterozygous carriers of a *PKLR* variant will also often have low PK activity; thus,
18 genetic testing has an important role as a confirmatory test (Bianchi, *et al* 2019). In
19 addition, genetic testing will often be required to establish the diagnosis in patients with
20 recent transfusions as well as those with suspected PKD and a normal PK enzyme
21 activity and PK:HK ratio. The second major conclusion is that erythrocyte PK enzyme
22 activity does not correlate with markers of clinical severity in PKD, but that erythrocyte
23 PK-R protein and ATP levels do, in particular the need for transfusions.

24 While a positive correlation between erythrocyte ATP level and 2,3-DPG level
25 and a negative correlation between erythrocyte ATP level and disease severity seem
26 counterintuitive, these findings may be due to the higher reticulocyte count noted in
27 patients with more robust hemolysis and lower hemoglobin levels (Figure 1C).
28 Reticulocytes have higher ATP on a per-cell basis than mature erythrocytes (Mentzer,
29 *et al* 1971), and higher reticulocyte counts in patients with severe disease are likely
30 indicative of the elevated erythropoietic drive necessary to compensate for more severe
31 hemolysis. PK-R protein levels appear to correlate with transfusion needs. Given that

1 the number of *PKLR* variants and genotype combinations limits predicting outcomes
2 based on molecular findings, erythrocyte PK-R protein levels may represent a better
3 marker of disease severity. Additional studies to further elucidate the prognostic value
4 of erythrocyte PK-R protein in patients could help to confirm whether this is a useful
5 future biomarker for clinicians managing patients with PKD.

6
7 **Acknowledgements:** H. Al-Samkari is the recipient of the National Hemophilia
8 Foundation-Shire Clinical Fellowship Award, the Harvard KL2/Catalyst Medical
9 Research Investigator Training Award, and the American Society of Hematology
10 Scholar Award.

11
12 **Authorship Contributions:** H. Al-Samkari wrote the first draft of the manuscript and
13 contributed to data analysis, creation of tables and figures, critical revision of the
14 manuscript, and final approval; K. Addonizio contributed to data collection, data
15 analysis, creation of tables and figures, critical revision of the manuscript, and final
16 approval; B. Glader provided red cell enzyme data and contributed to data collection,
17 critical revision of the manuscript, and final approval; D. Morton contributed to data
18 collection, critical revision of the manuscript, and final approval; S. Chonat contributed
19 to data collection, critical revision of the manuscript, and final approval; A. Thompson
20 contributed to data collection, critical revision of the manuscript, and final approval; K.
21 Kuo contributed to data collection, critical revision of the manuscript, and final approval;
22 Y. Ravindranath contributed to data collection, critical revision of the manuscript, and
23 final approval; H. Weng contributed to data collection, critical revision of the manuscript,
24 and final approval; J. Rothman contributed to data collection, critical revision of the
25 manuscript, and final approval; J. Kwiatkowski contributed to data collection, critical
26 revision of the manuscript, and final approval; C. Kung contributed to critical revision of
27 the manuscript and final approval; P. Kosinski contributed to data analysis, critical
28 revision of the manuscript and final approval; H. Al-Sayegh contributed to data analysis,
29 critical revision of the manuscript, and final approval; W. London contributed to data
30 analysis, critical revision of the manuscript, and final approval; R. Grace contributed to

1 study design, data collection, data analysis, creation of tables and figures, critical
2 revision of the manuscript, and final approval.

3

4 **Disclosures:** Al-Samkari: Consultancy (AgiOS, Dova, Moderna), Research funding
5 (AgiOS, Dova, Amgen). Addonizio: None. Glader: Scientific Advisor and Consultancy
6 (AgiOS), Research Funding (AgiOS). Morton: None. Chonat: Advisory board (Alexion,
7 AgiOS). Thompson: Consultancy (Bluebirdbio, Celgene, Novartis), Research Funding
8 (Bluebirdbio, Celgene, Novartis, Baxalta). Kuo: Honoraria (Alexion, Novartis);
9 Consultancy (AgiOS, Alexion, Bluebirdbio, Celgene, Novartis, Pfizer); Data Safety
10 Monitoring Board Chair (Bioverativ). Ravindranath: Research Funding (AgiOS). Weng:
11 None. Rothman: Research Funding (AgiOS, Novartis, Pfizer). Consultancy (AgiOS,
12 Novartis, Pfizer). Kwiatkowski: Consultancy (Celgene, Bluebirdbio, Imara, AgiOS).
13 Research Funding (Apopharma, Novartis, Terumo, Bluebirdbio). Kung: Employee
14 (AgiOS), Shareholder (AgiOS). Kosinski: Employee (AgiOS), Shareholder (AgiOS). Al-
15 Sayegh: None. London: Consultancy (United Therapeutics, ArQule, Inc.). Grace:
16 Scientific Advisor (AgiOS), Consultancy (AgiOS, Dova), and Research Funding (AgiOS,
17 Novartis).

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Table. Patient characteristics and results of erythrocyte metabolite and PK enzyme assays. RBC, red blood cell; DPG, diphosphoglycerate; ATP, adenosine triphosphate; PK-R, pyruvate kinase-red cell; PK, pyruvate kinase; HK, hexokinase; EU, enzyme unit; Hb, hemoglobin.

Patient Characteristics (N=41)	Value
Median age at enrollment (years, range)	25.3 (1.4-60.4)
Female	21 (51%)
Amish	21 (51%)
Splenectomy	33 (80%)
Total lifetime RBC transfusions, median (range) (N=34)	17 (0-149)
Red Cell Metabolite Levels	Result
2,3-DPG ($\mu\text{g/ml}$) level, median (range) (N=39)	796 (573-995)
Normalized 2,3-DPG level, median (range) (N=38)	0.0083 (0.0059-0.0126)
ATP ($\mu\text{g/ml}$) level, median (range) (N=39)	179 (78-233)
Normalized ATP level, median (range) (N=38)	0.0021 (0.0008-0.0029)
PK-R Protein Levels	Result
PK-R protein percentage ^a , median (range) (N=37)	8.70 (0.41-73.95)
Non-Amish, median (range) (N=16)	3.20 (0.41-73.95)
Missense/Missense, n=7	7.66 (0.41-73.95)
Missense/Non-missense, n=8	2.03 (0.72-51.71)
Non-missense/Non-missense, n=1	2.02 (-)
Amish (range), median (N=21)	9.06 (4.53-15.62)
Enzyme Activity (N=41)	Result
PK activity (EU/gm Hb), median (range)	1.1 (0.2-4.6)
HK activity (EU/gm Hb), median (range)	0.7 (0.3-1.4)
PK activity (category)	

Low (<3.2 EU/gm Hb)	37 (90%)
Normal (3.2-6.5 EU/gm Hb)	4 (10%)
High (>6.5 EU/gm Hb)	0 (0%)
PK:HK ratio ^b median (range)	1.88 (0.30-9.20)
Low PK/HK ratio	40 (98%)

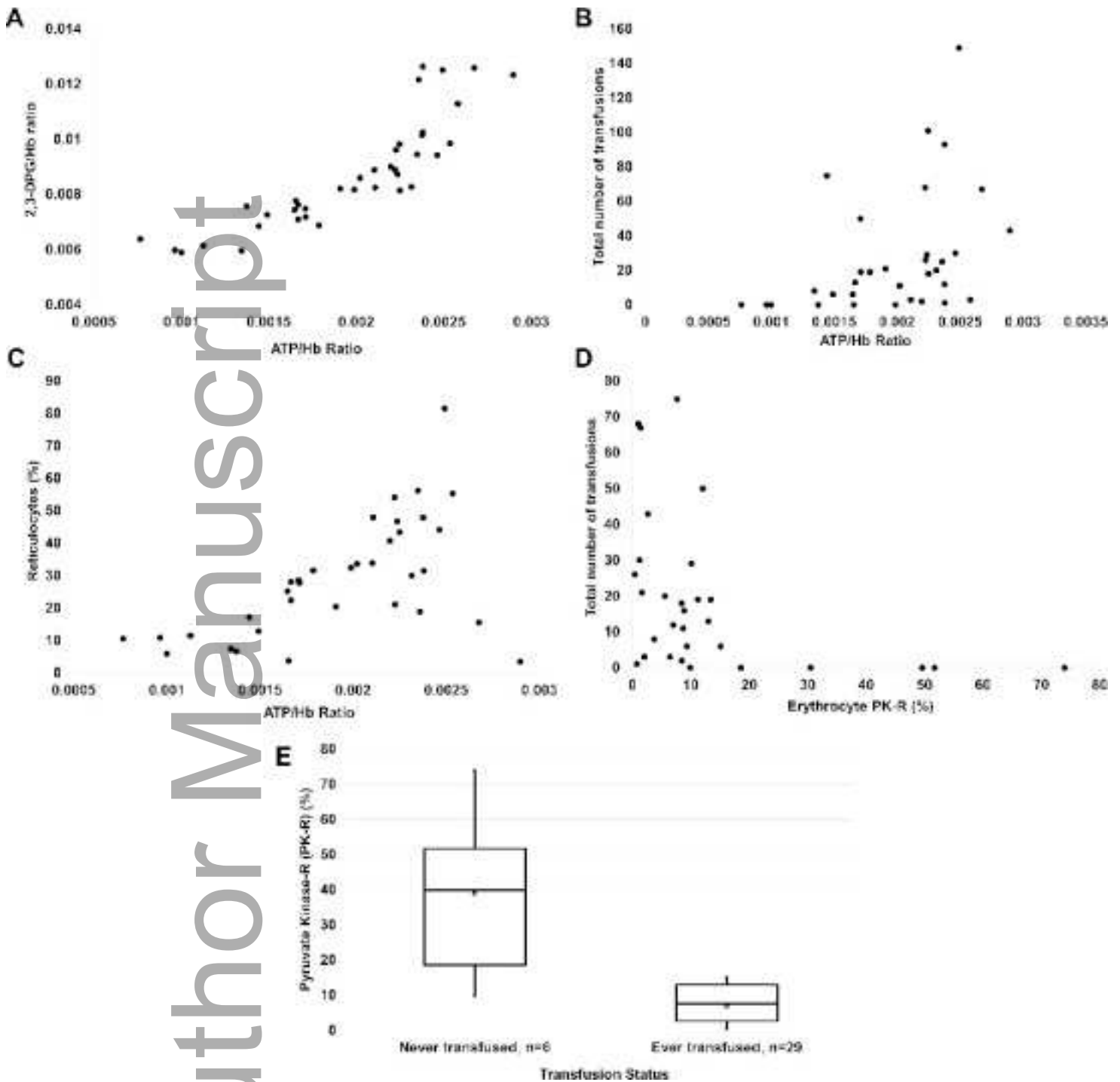
^aNormalized to hemoglobin concentration and expressed as the percentage of protein present in a healthy subject without PKD.

^bNormal range for PK:HK ratio is 8.7-22.5.

Figure 1. Relation of erythrocyte metabolites with one another or markers of clinical severity in PKD. (A) Erythrocyte ATP versus erythrocyte 2,3-DPG (N=38, $r=-0.93$; $P<0.0001$). (B) Erythrocyte ATP versus total number of lifetime RBC transfusions (N=33, $r=0.527$, $P=0.0016$). (C) Erythrocyte ATP versus percent reticulocyte count (N=35, $r=0.581$, $P=0.0003$). (D) Erythrocyte PK-R protein versus total number of lifetime RBC transfusions (N=30, $r=-0.527$, $P=0.0028$). (E) Erythrocyte PK-R protein level in never transfused (N=6) versus ever transfused (i.e., received at least one red cell transfusion over the lifetime, N=29) patients ($P=0.0014$).

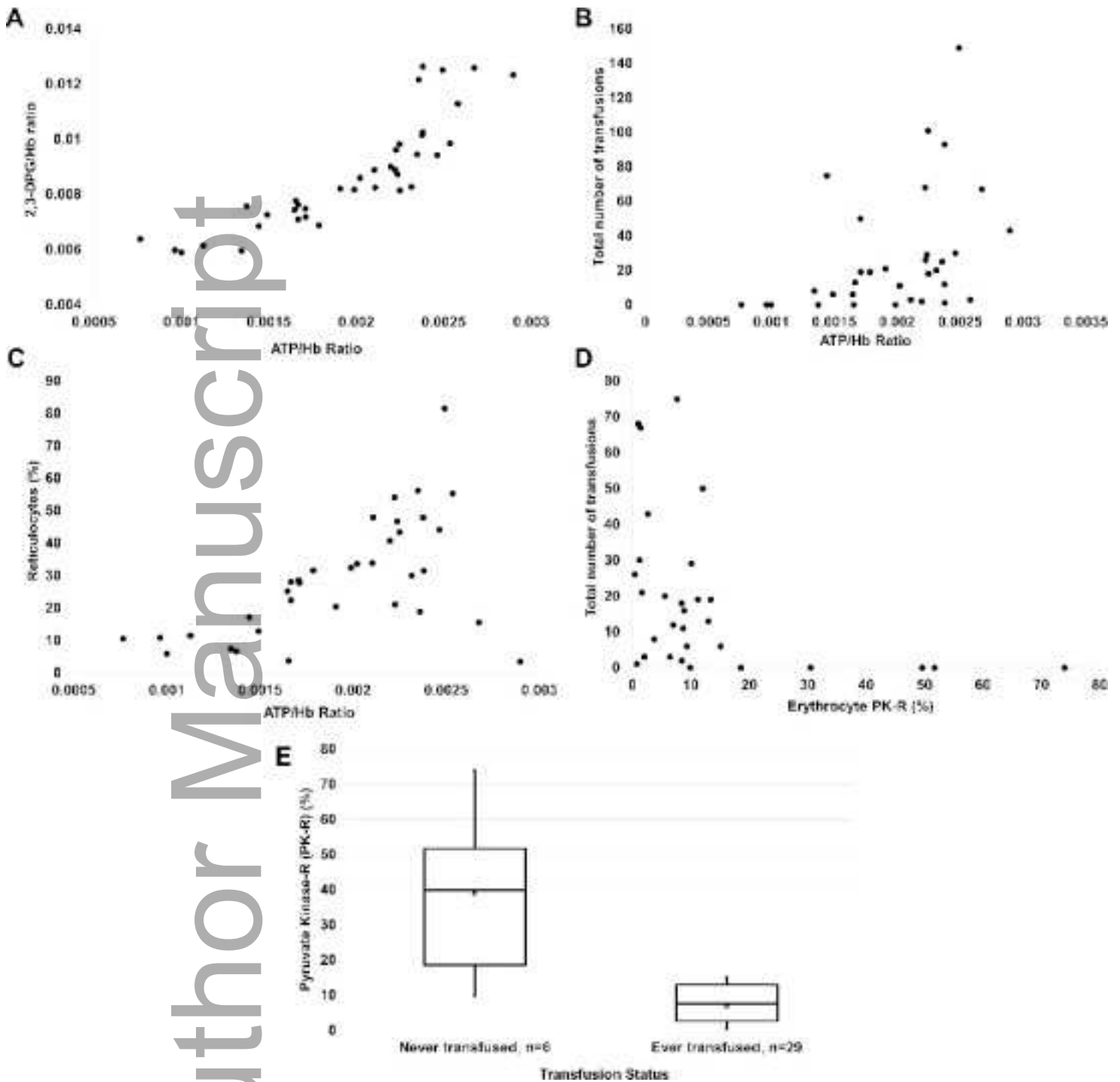
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