



Utility of the succinate:fumarate ratio for assessing SDH dysfunction in different tumor types[☆]



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ARTICLE INFO

Article history:

Received 17 October 2016

Accepted 17 December 2016

Available online 30 December 2016

Keywords:

Succinate dehydrogenase deficiency

Mass spectrometry

Pheochromocytoma and paraganglioma

Gastrointestinal stromal tumor

Renal cell carcinoma

ABSTRACT

Objective: Mutations of genes encoding the four subunits of succinate dehydrogenase (SDH) have been associated with pheochromocytoma and paraganglioma (PPGLs), gastrointestinal stromal tumors (GISTs) and renal cell carcinomas (RCCs). These tumors have not been characterized in a way that reflects severity of SDH dysfunction. Mass spectrometric analysis now allows measurement of metabolites extracted from formalin fixed paraffin embedded (FFPE) specimens. We assess whether SDH deficiency in various tumor types characterized by loss of SDHB protein expression correlates with SDH dysfunction as assessed by the ratio of succinate:fumarate in FFPE specimens.

Patients and methods: Sections of FFPE tumor specimens from 18 PPGL, 10 GIST and 11 RCC patients with known *SDHx* mutation status for SDH deficiency were collected for mass spectrometric analysis of succinate and fumarate.

Results: FFPE samples showed higher succinate:fumarate ratios in SDH-deficient PPGLs compared to SDH-sufficient PPGLs. Similarly, a higher succinate:fumarate ratio was able to distinguish SDH-deficient GISTs and RCCs from their SDH-sufficient counterparts with great selectivity. Interestingly, the cut-off value of the succinate:fumarate ratio was two-folds greater in RCCs than GISTs.

Conclusion: Analyzing biochemical imbalances preserved in FFPE specimens with mass spectrometry expands the method and sample type repertoire available for characterisation of multiple neoplasias associated with SDH deficiency.

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1. Introduction

Deregulation of cellular metabolism has been widely recognized as one of the hallmarks of cancer [1]. Accumulation of biomolecules as a result of enzyme impairment has gained interest in the field of cancer biomarker discovery [2]. Mutations in genes encoding subunits of succinate dehydrogenase (SDH) leading to loss or dysfunction of the mitochondrial enzyme involved in both the Krebs cycle and electron transport chain have been associated with multiple neoplasias [3]. These include pheochromocytomas and paragangliomas (PPGLs), gastrointestinal stromal

tumors (GISTs) and renal cell carcinomas (RCCs) [4–6]. Clinical presentations, immunohistochemical (IHC) staining of SDH subunit proteins SDHA/B from resected specimens, and genetic screening of *SDHx* genes are typically used to characterize these tumors [7,8]. SDH-deficient tumors, characterized by loss of SDHB IHC, accumulate succinate and secondarily hypoxia inducible factors (HIFs) with downstream consequences referred to as pseudohypoxia [9]. Advances in liquid chromatography tandem mass-spectrometry (LC-MS/MS) allow sensitive measurement of accumulated succinate due to SDH dysfunction [10]. Metabolomic profiling of fresh frozen PPGL tumors suggested a direct link between tumor-associated *SDHx* mutations and the measurable metabolite imbalance of a high succinate:fumarate ratio [11,12].

Fresh frozen specimens represent the gold standard sample type for metabolomic analyses, however storage and availability varies greatly across different sites limiting their use [13]. Formalin-fixed paraffin embedded (FFPE) specimens constitute a widely available and accessible archived pathological resource representing a good alternative for

[☆] This work was supported by the Hillcrest Foundation (Perpetual Trustees) (IPAP201600914) and by NHMRC project grant 1108032.

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fresh-frozen tissue especially in studies involving rare cancers [14]. The use of FFPE tissues in retrospective metabolomic studies has shown potential applicability for LC-MS/MS based profiling of sarcomas and thyroid cancer tissue [15–17]. This led us to question whether FFPE specimens from SDH-deficient tumors could be used for metabolomic profiling.

The aim of this study was to assess whether elevated succinate:fumarate ratios, as assessed by LC-MS/MS from FFPE samples, are a consistent biochemical signature in SDH-deficient GISTs and RCCs in the same manner as previously observed in PPGLs.

2. Material and methods

2.1. FFPE specimens

FFPE specimens from 18 cases of PPGLs encompassing 9 pheochromocytoma and 9 paraganglioma patients, 10 cases of GISTs and 11 cases of RCCs were obtained from surgical resection of primary tumors. The use of specimens in this study was approved by Northern Sydney Local Health District Human Research Ethics Committee (LNR 1312-417M). The presence of neoplastic tissue in the sections were confirmed by an experienced surgical pathologist (AJG) in all samples and all samples were confirmed to have a neoplastic cellularity of >60%. SDH-deficiency was defined by the standard IHC approach requiring loss of SDHB protein expression in neoplastic cells, with preserved expression in non-neoplastic cells which act as an internal positive control [18], whereas SDH-sufficient specimens were defined by positive granular cytoplasmic SDHB staining. *SDHx* mutation status was confirmed in SDH-deficient specimens from PPGL group with 2 *SDHA*, 5 *SDHB*, 2 *SDHD* germline mutations and RCC group with 2 *SDHB*, 1 *SDHD* germline mutations. In GIST group, *SDHA* mutation was confirmed in 2 of 5 SDH-deficient tumors (Supplementary Table 1). However, based on previous reports, approximately half of SDH-deficient GISTs have no underlying *SDHx* mutation [6,18,19]. The first section cut from each FFPE specimen block was discarded to minimize contamination due to prolonged environmental exposure. Two 50 μm sections were then cut from each block and pooled into a 1.5 mL microfuge tube [12].

2.2. Sample preparation

Succinate and fumarate certified reference materials were purchased from Sigma-Aldrich (Australia) for the creation of calibration curves. $^{13}\text{C}_4$ succinate was also purchased from Sigma-Aldrich and $^{13}\text{C}_4$ fumarate was purchased from Cambridge Isotope Laboratories (USA) for use as internal standards. 1 mL of extraction solution (20% LC/MS grade H_2O , 80% LC/MS grade methanol containing 0.1 $\mu\text{g}/\text{mL}$ $^{13}\text{C}_4$ fumarate and 0.5 $\mu\text{g}/\text{mL}$ $^{13}\text{C}_4$ succinate) was added to each microfuge tube containing FFPE sections. After a brief vortex, mixtures were incubated for 45 min at 70 $^\circ\text{C}$ followed by 5 min on ice. Metabolites were extracted from melted paraffin by collecting and combining supernatants from two consecutive centrifugations at 14,000 $\times g$ for 10 min at 4 $^\circ\text{C}$. Samples were dried by evaporating methanol for 2 h using a Speedvac and then stored at -80°C until further use [12].

2.3. LC-MS/MS analysis

Analysis of extracts was performed on a Prominence high performance liquid chromatography system (Shimadzu, Australia) coupled to an API QTRAP 5500 mass spectrometer (SCIEX, Australia) operated in negative electrospray ionisation. Separation of target analytes from isobaric interferences was achieved using an Ascentis Express 100 \times 3.0 mm 2.7 μm RP Amide (Sigma Aldrich, Australia) analytical column held at 40 $^\circ\text{C}$ and isocratic elution using aqueous 0.4% formic acid with a flow rate of 0.5 mL/min. Succinate eluted at 1.57 min and fumarate eluted at 2.29 min with chromatographic resolution between interferences and a total run time of 3 min (Fig. 1).

Initially 1 μL of extracted samples were injected onto the system with a linear calibration range of 22.5–90,000 ng/mL for succinate and 9–450 ng/mL for fumarate. Extracts that demonstrated fumarate levels between the lower two calibrators (0.9 and 9 ng/mL) were reinjected, along with calibrators, with a 10 μL injection for quantitation to 0.9 ng/mL. At the end of every batch the column was cleaned with acetonitrile at 0.8 mL/min for 10 min to remove any strongly retaining compounds. Analytes were detected by monitoring multiple reaction monitoring m/z transitions of 116.9 > 73.1 and 116.9 > 99.1 as quantifier and qualifier transitions, respectively, for succinate, 121.0 > 76.1 for the $^{13}\text{C}_4$ succinate internal standard, 114.9 > 71.0 for fumarate and 119 > 74 for the $^{13}\text{C}_4$ fumarate internal standard. Unlike previous reports we were unable to find a suitable qualifier transition for fumarate.

2.4. Statistical analysis

Statistical analyses of LC-MS/MS results were performed using Prism 6.0 f. Unpaired two-tailed t -tests were used to determine the differences in succinate:fumarate ratio between SDH-deficient and SDH-sufficient tumor groups. Results were considered significant if $p < 0.05$. The optimal cut-off value for discriminating SDH-deficient tumors using the succinate:fumarate ratio was determined by generating a receiver-operating characteristic (ROC) curve. Values are represented as mean \pm SEM unless otherwise stated.

3. Results

3.1. SDH-deficient vs SDH-sufficient PPGLs

Analysis of metabolites extracted from FFPE PPGL specimens showed a significantly higher succinate:fumarate ratio in SDH-deficient PPGL tumors (185.75 ± 57.24 , $n = 9$) when compared to SDH-sufficient PPGL tumors (40.02 ± 12.11 , $n = 9$) ($p < 0.05$) (Fig. 2). Although statistical analysis of these two groups showed a significantly different metabolite imbalance, a cut-off ratio of succinate:fumarate (63.12) generated by ROC curve (Supplementary Fig. 1) could not clearly separate these tumor types; 1 false positive and 2 false negatives (Supplementary Table 2). Interestingly among SDH-sufficient samples, those with highest succinate:fumarate ratios were from specimens with germline *VHL* mutations. The succinate:fumarate ratio was not able to distinguish *VHL* mutated PPGLs from SDH-deficient PPGLs (Supplementary Fig. 2).

3.2. SDH-deficient vs SDH-sufficient GISTs

The succinate:fumarate ratio in five SDH-deficient GISTs (73.25 ± 22.06) was shown to be significantly higher than five SDH-sufficient GISTs (representing patients with *KIT* or *PDGRA* gene mutations; 15.84 ± 2.65) ($p < 0.05$) (Fig. 3). Using logistic regression, an area under the ROC curve of 1.00 was calculated and a cut-off succinate:fumarate ratio of 23.48 was determined that distinguished SDH-deficient and SDH-sufficient GISTs (Supplementary Fig. 3).

3.3. SDH-deficient vs SDH-sufficient RCCs

The mean succinate:fumarate ratio in three SDH-deficient RCCs (46.9 ± 2.234) was significantly higher than eight SDH-sufficient RCCs (19.30 ± 4.53) ($p < 0.05$) (Fig. 4). With an area under the ROC curve of 1.00, a succinate:fumarate ratio of 43.79 was calculated as the cut-off value that distinguished SDH-deficient RCCs from SDH-sufficient RCCs (Supplementary Fig. 4).

4. Discussion

Expression of functional SDH in the mitochondria is essential for cellular respiration and energy production [20,21]. “Pseudohypoxia”, in

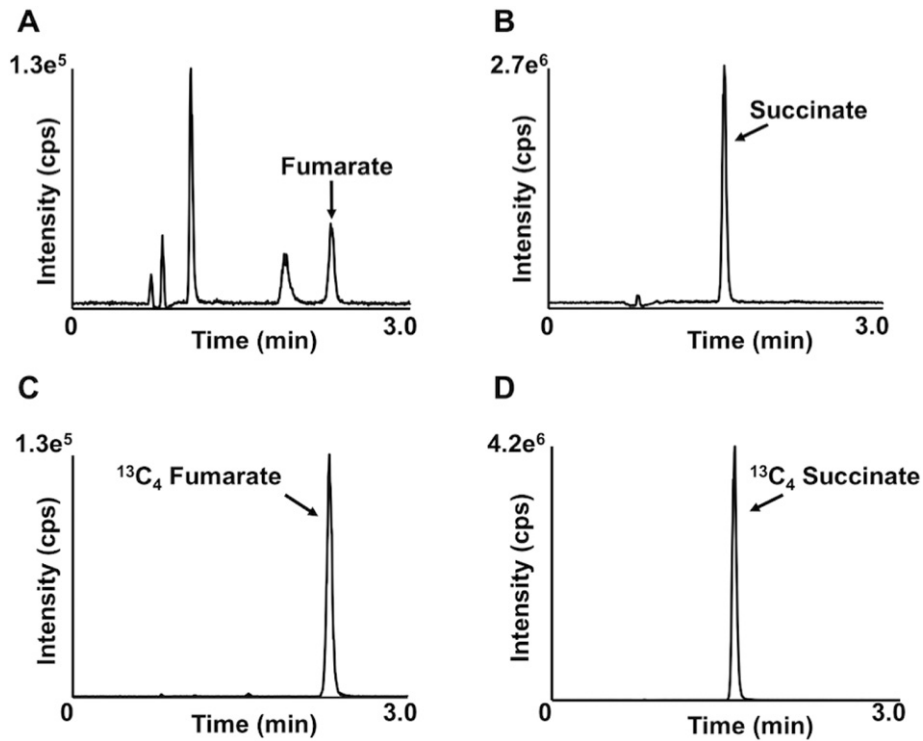


Fig. 1. Multiple reaction monitoring chromatograms from an extracted FFPE sample for (A) fumarate (B) succinate (C) $^{13}\text{C}_4$ fumarate and (D) $^{13}\text{C}_4$ succinate.

which HIF is stabilized despite normoxic conditions as a consequence of succinate-mediated prolyl hydroxylase domain (PHD) inhibition, has been associated with SDH-deficient neoplasias such as PPGLs, GISTs and RCCs [22,23]. Analyzing the substrate-to-product ratio provides direct assessment of enzymatic function and this has been implemented in the discovery of accumulated metabolites associated with cancer, now labelled as ‘onco-metabolites’ [2]. High succinate measurement relative to fumarate in the tumors represents a direct link to functional aspects associated with SDH-deficiency. In this study we have shown for the first time that an elevated succinate:fumarate ratio is a consistent biomolecular phenotype of SDH-deficient tumors that include GISTs and RCCs as well as PPGLs. We confirmed a previous report that

measurement of these metabolites is possible from FFPE specimens [12]. Of note, one of our SDH-deficient specimens had been stored for 25 years in archive.

Our study is the first to report significantly higher succinate:fumarate ratios in SDH-deficient GISTs and RCCs compared to corresponding SDH-sufficient GISTs and RCCs. Clear separation of GIST and RCC tumor groups with 100% specificity suggests potential diagnostic application of LC/MS-MS based profiling of metabolites extracted from FFPE specimens as previously observed for PPGLs. Interestingly, our cut-off succinate:fumarate ratio determined by ROC curve analysis was almost two-fold higher for discriminating SDH-deficient RCCs compared to SDH-deficient GISTs. This suggests the

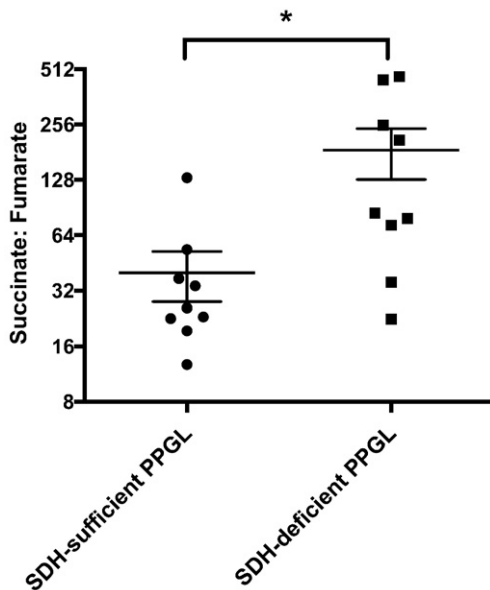


Fig. 2. Scatter plot comparing succinate:fumarate ratio between SDH-sufficient (n = 5) and SDH-deficient PPGLs (n = 9), error bars representing SEM from mean (*p < 0.05).

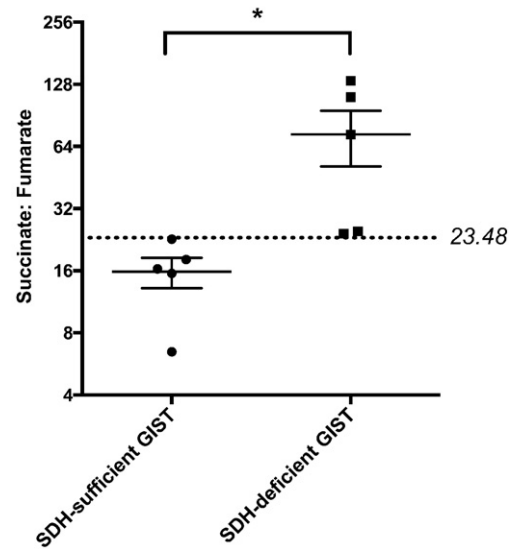


Fig. 3. Scatter plot comparing succinate:fumarate ratio between SDH-sufficient (n = 5) and SDH-deficient GISTs (n = 5), error bars representing SEM from mean (*p < 0.05). ROC curve analysis was used to determine cut-off value of 23.48 marked as a dashed line.

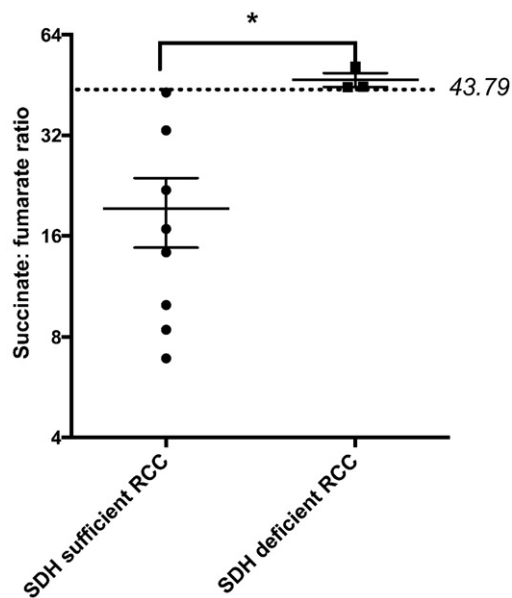


Fig. 4. Scatter plot comparing succinate:fumarate ratio between SDH-sufficient ($n = 8$) and SDH-deficient RCCs ($n = 3$), error bars representing SEM from mean ($*p < 0.05$). ROC curve analysis was used to determine cut-off value of 43.79 marked as a dashed line.

extent of biochemical imbalance caused by SDH-deficiency is likely to be tissue-specific.

We have confirmed the earlier finding by Richter et al. [12] that *VHL*-mutated PPGLs have higher succinate:fumarate ratios within the SDH-sufficient group (Supplementary Fig. 2), suggesting that *VHL*-deficiency causes functional SDH-deficiency. PPGL tumor specimens with *VHL* gene mutations have been associated with variably impaired SDHB protein expression [24], and although the mechanism for this is unknown several hypotheses have been advanced including pseudohypoxic expression of miR-210 that in turns downregulates SDHD expression [25].

Just as higher succinate levels were detected in nuclear magnetic resonance (NMR) spectroscopy of *SDHx* tumors in vivo [26,27], direct measurement of succinate and fumarate by LC-MS/MS based analysis appears therefore to be a powerful technique to interrogate SDH function ex vivo. These methods now allow quantitative analysis of SDH dysfunction in a manner that complements qualitative assessment of SDHB protein expression by IHC. Elevated succinate:fumarate appears to have at least three potential causes: (a) inactivation by germline mutation in *SDHx* gene (together with somatic loss of the other allele) herein shown to be true for *SDHx*-mutated RCCs as well for PPGLs; (b) hypermethylation of the *SDHC* promoter, as first described in Carney triad [28] and more recently in PPGL by Richter et al. [29], and which is possibly also the mechanism for SDH-deficiency in at least some of our GIST samples; and (c) by germline *VHL* mutation which is associated with variably impaired SDH function.

Our data also sound a note of caution in that a very few *SDHx*-mutated PPGLs were associated with normal succinate:fumarate values (i.e. 'false negatives', Supplementary Table 2). The *SDHx*-mutated PPGL with the lowest succinate:fumarate (i.e. indistinguishable from SDH-sufficient samples) contained *SDHB* mutation c.380T>G, p.Ile127Ser. This was particularly interesting to us, since we have recently described a method for measuring mutant SDH function in vitro, and this same mutation was found to have minimal impact on enzymatic function [30]. It is intriguing therefore to speculate that SDH-deficient PPGLs with normal succinate:fumarate ratios are associated with mutations that do not primarily alter SDH activity and may otherwise cause tumor development through alternative mechanisms (e.g. via electron transport abnormalities).

A limitation of this study was the heterogeneous nature and variable tumor content in sections of our samples. Although all of our samples were reviewed by a pathologist to confirm the presence of neoplastic tissue in relatively high cellularity (>60% neoplastic in all cases), we did not specifically micro-dissect the sections, and we acknowledge that there may have been dilution of the metabolite signal by normal cells in the surrounding tissue. However, despite this limitation, a significant difference in the succinate:fumarate ratio between the SDH-sufficient and -deficient group was still observed in most of the samples and tumors with similar morphologies and growth patterns could be expected to demonstrate similar neoplastic cellularities. That is, SDH-deficient GISTs would have similar neoplastic cellularity to other GISTs, and SDH-deficient PPGLs could be expected to have similar neoplastic cellularity to other PPGLs and so on.

5. Conclusion

We have demonstrated that SDH-deficient PPGLs, GISTs and RCCs retain a significantly higher succinate:fumarate ratio compared to SDH-sufficient counterparts, and can be readily assayed from archived samples. Our study highlights the importance of metabolomic analysis to directly measure SDH dysfunction that is associated with multiple types of neoplasias.

Supplementary data to this article can be found online at <http://dx.doi.org/10.1016/j.ymgmr.2016.12.006>.

Disclosure statement

The authors have nothing to disclose.

Acknowledgments

This work was supported by the Hillcrest Foundation (Perpetual Trustees) to DB, and by NHMRC project grant 1108032 to RCB, TD, DB and AG.

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