Science

February 2005, Volume 307 (5712), Pages 1088-1091 http://dx.doi.org/10.1126/science.1105692
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Iron Isotope Constraints on the Archean and Paleoproterozoic Ocean Redox State

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Abstract:

The response of the ocean redox state to the rise of atmospheric oxygen about 2.3 billion years ago (Ga) is a matter of controversy. Here we provide iron isotope evidence that the change in the ocean iron cycle occurred at the same time as the change in the atmospheric redox state. Variable and negative iron isotope values in pyrites older than about 2.3 Ga suggest that an iron-rich global ocean was strongly affected by the deposition of iron oxides. Between 2.3 and 1.8 Ga, positive iron isotope values of pyrite likely reflect an increase in the precipitation of iron sulfides relative to iron oxides in a redox stratified ocean.

The rise of atmospheric oxygen, which began by about 2.3 Ga (1-3), was one of the most important changes in Earth's history. Because Fe, along with C and S, are linked to and maintain the redox state of the surface environment, the concentration and isotopic composition of Fe in seawater were likely affected by the change in the redox state of the atmosphere. The rise of atmospheric oxygen should have also led to dramatic changes in the ocean Fe cycle because of the high reactivity of Fe with oxygen. However, deposition of banded iron formations (BIFs) during the Paleoproterozoic era suggests that the deep ocean remained anoxic, at least episodically, until about 1.8 Ga, which allowed high concentrations of Fe(II) to accumulate in the deep waters (4).

Here we use Fe isotope systematics ($\underline{5}$) to provide constraints on the redox state of the Archean and Paleoproterozoic oceans and to identify direct links between the oxidation of the atmosphere and the Fe ocean cycle. Laboratory and field studies suggest that Fe isotope variations are associated mainly with redox changes ($\underline{6}$, $\underline{7}$). Lithogenic sources of Fe on the modern oxygenated Earth, such as weathering products, continental sediments, river loads, and marine sediments, have isotopic compositions similar to those of igneous rocks ($\underline{8}$, $\underline{9}$). In contrast,

seafloor hydrothermal sulfides and secondary Fe-bearing minerals from the altered oceanic crust span nearly the entire measured range of δ^{56} Fe (5) on Earth from -2.1 to 1.3‰ (10, 11). Large variations of δ^{56} Fe (from -2.5 to 1.0‰) in Late Archean to Early Paleoproterozoic BIF have been also reported (12) highlighting the roles of ferrous Fe oxidation, fluid-mineral isotope fractionation, and potentially microbial processes in the fractionation of Fe isotopes.

Study of S isotope composition of sedimentary pyrite over geological time has placed important constraints on the S cycle and the evolution of ocean chemistry (13) and here we apply a similar time-record approach to explore potential change in Fe isotope compositions. Pyrite formation in modern organic-rich marine sediments is mediated by sulfate-reducing bacteria and proceeds essentially through the dissolution and reduction of lithogenic Fe-oxides and Fesilicates to Fe(II), either below the sediment-water interface or in stratified euxinic bottom waters (14-16). During reduction of Fe-oxides, diagenetic fluids with isotopically light Fe(II) may be produced (17, 18). However, the Fe isotope composition of sedimentary pyrite from Phanerozoic organic-rich sediments studied so far (Fig.1, Table S2, 19) suggests that such processes are unlikely to produce sedimentary pyrite with δ^{56} Fe < -0.5‰. Presumably, most of reactive Fe is scavenged to form pyrite, minimizing Fe isotope fractionation regardless of the isotope effect during Fe reduction (17) and precipitation (20). In contrast, when high concentrations of Fe(II) accumulate under anoxic conditions and low sulfide concentration, large δ^{56} Fe variations (10-12) may occur due to partial Fe(II) oxidation, Fe(III) reduction and distillation processes during mineral precipitation. We thus hypothesize that Fe isotope variations in sedimentary pyrite are particularly sensitive to the concentration of dissolved Fe(II) and can be used to place important constraints on the sources and sinks of the Fe(II) reservoir.

We analyzed Fe isotope composition of sulfides in black shales ranging in age from Precambrian to Late Cretaceous, specifically focusing on Late Archean to early Paleoproterozoic time (Fig.1, 21). The emerged general pattern of Fe isotope record suggests that the Earth's history may be divided into three stages which are strikingly similar to the stages defined by the δ^{34} S, Δ^{33} S, and δ^{13} C records as well as other indicators of the redox state of the atmosphere and ocean (2, 3, 13, 22).

Stage I extends from before 2.8 Ga to ca. 2.3 Ga and is characterized by highly variable and negative δ^{56} Fe values. The entire range of δ^{56} Fe between 0.5 to -3.5% is often observed within a single section of black shales but individual pyrite nodules from the same stratigraphic level have similar δ^{56} Fe (Fig.2). Since dissimilatory Fe(III) reduction has been suggested to be important on the early Earth (23) and is known to produce significant Fe isotope fractionations (6, 24), it can be hypothesized that these extreme Fe isotope fractionations were produced by this metabolic activity. Three independent observations argue against this hypothesis. First, Fe isotope fractionation during single-step bacterial reduction of Fe-oxides (with initial δ^{56} Fe at 0%) is unlikely to produce Fe(II) with δ^{56} Fe of less than -1.3% (17). Secondly, if δ^{56} Fe as low as -3.5% can be generated through multiple steps of Fe oxidation and reduction, then the evidence for these processes should be evident in younger sediments, but they are not documented (Fig.1). In addition, bacterial Fe(III) reduction is expected to produce pyrite with locally highly variable negative δ^{56} Fe depending on the extent of Fe(III) reduction and Fe(II) re-oxidation. Our samples (Fig.2, Table S2) do not show significant variability between individual sulfide nodules, and suggest a common source of Fe(II). Thirdly, the amount of biogenically produced Fe(II) would need to be unrealistically high during the Archean to swamp the global influence of hydrothermally derived Fe(II) with δ^{56} Fe between 0 and -0.5% (8) delivered to the deep ocean.

 δ^{56} Fe as low as -2.3% have been observed in Fe-rich groundwater springs precipitating isotopically heavy ferrihydrite along a fluid flow path (25) and yielding low δ^{56} Fe in residual Fe(II) pool. Adsorption of Fe(II) onto Fe oxide particles may also provide an additional mean to produce isotopically negative Fe(II) pool through the preferential sorption of ⁵⁶Fe onto Fe oxide surfaces (24). In a similar manner, low δ^{56} Fe for Archean oceans may reflect preferential sequestration of ⁵⁶Fe on Fe oxides (26, 27). Indeed magnetite and hematite in BIFs are often characterized by positive δ^{56} Fe (12), such as for example δ^{56} Fe up to 1.6% in ca. 2.7 Ga Iron Formations of the Belingwe Belt, Zimbabwe (Fig.1). Large stratigraphic variations in δ^{56} Fe of sedimentary pyrites in 2.7 to 2.5 Ga black shales, up to 3‰ over tens to hundreds meters of section (Fig.2), are suggestive of changes in Fe isotope composition of seawater over short periods of time on the order of few Ma. This implies non-steady state of the Archean Fe cycle with variable Fe concentrations due to the competitive effect of Fe-oxide precipitation and Fe supply from hydrothermal sources. These rapid changes of Fe concentrations are consistent with the proposition that Fe-oxide deposition in BIFs resulted from the episodic upwelling of Fe-rich deep waters accompanied by partial biological and/or abiological oxidation (26) in shallow waters (28). Alternatively, Fe-oxide deposition within marine sediments on continental shelves or in deep-ocean may have also provided an important sink for Fe between periods of large BIF deposition.

We used a simple Rayleigh distillation model to explore the influence of Fe-oxide deposition on Fe isotope composition of seawater (29). Fe is delivered to the ocean from rivers and seafloor hydrothermal systems with δ^{56} Fe ranging from 0.0 to -0.5‰ (8) and is removed by precipitation of Fe-oxides. δ^{56} Fe as low as -3.5‰ are only reached when more than 90% of Fe of the initial Fe pool is precipitated as Fe-oxides. δ^{56} Fe of -1.5‰ to -2.0‰, which are more typical

of Late Archean sulfides, correspond to about 50% of Fe precipitated as oxides. This value is similar to the estimates of Fe sink in BIFs based on P adsorption (30).

Stage II, covering the time interval from ca. 2.3 to ca. 1.7 Ga, is characterized by the disappearance of negative δ^{56} Fe and the emergence of positive δ^{56} Fe up to 1.2‰. Major perturbations in biogeochemical and climatic record occurred during the beginning of Stage II. These include a) negative and positive carbon isotope excursions in carbonates sandwiched between two glacial diamictites; b) Earth's earliest global glaciations; and c) oxidation of the Earth's atmosphere as suggested by increasing seawater sulfate content inferred from the δ^{34} S record and appearance of sulfate evaporites, disappearance of non-mass dependent S isotope fractionation, appearance of red beds, oxidized paleosols, hematitic oölites and pisolites, Mnoxide deposits, and Ce anomalies in chemical sedimentary deposits (3, 13, 22, 31). Strikingly, the appearance of positive δ^{56} Fe, persisting until ca 1.7 Ga, together with the disappearance of strongly negative δ^{56} Fe occur during the period when the most sensitive indicators for the rise of atmospheric oxygen first appear. All these observations suggest that the oxidation of the surface environment in the early Paleoproterozoic was relatively rapid and directly affected Fe isotope composition of the ocean.

How the change in the Fe isotope record ca. 2.3 Ga ago corresponds with change of oceanic Fe cycle and redox state of the ocean is not straightforward. Interestingly, large BIF deposits are almost entirely lacking between 2.3 and 2.1 Ga (32) which is consistent with the lack of negative δ^{56} Fe during this period. However, BIF deposition returned at ca. 2.1 Ga and major BIFs were deposited in North America and Australia (32). If δ^{56} Fe of pyrites in black shales deposited between 2.1 to 1.8 Ga are representative of the whole ocean, then BIF deposition mechanisms were different from those prevailing during the Archean. We infer that late

Paleoproterozoic BIFs were deposited in an oxygenated layer of the ocean and complete precipitation of Fe from Fe-rich plumes uplifting from the deep ocean did not affect Fe isotope composition of the deep ocean. Despite the limited number of analyses, the narrow range of δ^{56} Fe of hematite and magnetite from the 1.88 Ga Biwabik and Tyler formations compared to Archean BIFs (Fig. 1; Table S3) is consistent with this assumption.

An important consequence of the rise of atmospheric oxygen levels was the initiation of oxidative weathering and an increase in sulfate delivery to seawater (13). Consequently, the formation of Fe sulfides in the water column of pericratonic basins may have became the dominant part of the global ocean Fe cycle and may have prevented deposition of large BIFs, except during periods of intense submarine volcanic activity followed by high hydrothermal input of Fe. The effect of the increased role of sulfide production on the Fe isotope record is presently uncertain since reliable estimates of equilibrium Fe isotope fractionation during pyrite formation are lacking (20). One plausible hypothesis is that the positive δ^{56} Fe in 2.3 Ga to 1.8 Ga sedimentary sulfides might be related to sulfide precipitation from a Fe-rich pool with δ^{56} Fe composition around 0% and a pyrite-Fe(II) fractionation factor of up to 1% as suggested in previous studies (10, 33). This indicates that sulfide produced by sulfate reducing bacteria during this period has been completely titrated by dissolved Fe species in the Fe-rich and sulfide-poor ocean.

The disappearance of major BIFs after ca. 1.8 Ga is thought to indicate that the deep ocean became either progressively oxic or euxinic (4). Since the solubility of Fe-sulfides and Fe-oxides is low, most of hydrothermally-derived Fe(II) was likely rapidly precipitated in the deep ocean allowing few possibilities to produce Fe isotope fractionation. The lack of significant δ^{56} Fe variations in sulfides from black shales younger than 1.5 Ga is thus consistent with the general

picture that the whole ocean was Fe-poor after ca. 1.6 Ga but do not place constraints on the oxic *vs.* sulfidic nature of the deep ocean. Evidently, more data covering the Phanerozoic, Meso- and Neoproterozoic time intervals are required to fully understand the change in ocean Fe cycle at ca. 1.8 Ga.

Our Fe isotope record provides new insights into the Archean and Paleoproterozoic ocean chemistry and redox state. Fe isotopes suggest that the Archean oceans were globally Fe-rich, their Fe isotope composition and Fe content were variable in response to the episodic establishment of an Fe-rich pool supplied by hydrothermal activity and deposition of Fe-oxides, either in BIFs or dispersed throughout sediments on continental shelves and in the deep sea. After the rise of atmospheric oxygen by ca. 2.3 Ga, Paleoproterozoic ocean became stratified and characterized by an increase of sulfide precipitation relative to Fe-oxide precipitation. During this period, BIFs were likely deposited by upwelling of Fe(II)-rich plumes and rapid oxidation in the oxygenated layer of the ocean. Conducting Fe isotope analyses of sedimentary sulfides in conjunction with S isotope analyses should enable a more refined understanding of the origin of the positive Fe isotope excursion and the biogeochemical cycles of Fe and S during the Paleoproterozoic.

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- 5. Data are reported using the delta notation relative to IRMM-14 Fe isotope reference standard defined as δ⁵⁶Fe=1000 * ((⁵⁶Fe/⁵⁴Fe) _{sample} / (⁵⁶Fe/⁵⁴Fe) _{IRMM-14} 1). External precision of δ⁵⁶Fe are estimated at 0.10‰ (2σ level). Analytical procedures, sample descriptions and Fe isotope composition of various georeference materials, black shales, and BIF are available as supporting materials on *Science Online*.
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- 21. Samples from different black shale units of similar ages were selected to distinguish between local and global effects on Fe isotope composition of seawater. We extracted between 10 to 20 sulfide grains to obtain a best estimate for the average δ^{56} Fe value for each sample. For

- comparison and to evaluate heterogeneity, we also analyzed several individual sulfide nodules from the same samples (Table S2).
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- 27. Magnetite-Fe(II) fractionation factor of approximately 2.4‰ has been inferred from BIFs data (12), which is slightly less than the equilibrium Fe(III)-Fe(II) fractionation factor of 2.9‰ at 22°C (42). The fractionation between ferrihydrite and Fe(II) of 1.5‰ measured during anaerobic photosynthetic Fe(II) oxidation by bacteria (43) is slightly larger than the 0.9‰ fractionation measured during abiotic oxidation of Fe(II) to ferrihydrite (25).
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$$\delta^{56}Fe = (1000 + \delta^{56}Fe_i) * F^{(\alpha BIF\text{-}1)}$$
 -1000

- where δ^{56} Fe_i is the initial value of hydrothermally derived Fe(II) at -0.5% (8); α_{BIF} is the fractionation factor during Fe oxidation and Fe oxide precipitation, ranging between 1.0015 and 1.0023 (6).
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Petsch, and H. Baioumy for advise and access to sample collections and Lary Ball for technical assistance. OJR is grateful to M. Bickle and A. Galy from the University of Cambridge for the access to Belingwe Iron Formation samples and analytical support of the Nu Plasma. AB fieldwork in South Africa and Western Australia was supported by NASA and PRF grants to H.D. Holland and by ARC and MERIWA grants to M. Barley and B. Krapez. We thank J.M. Hayes, H.D. Holland and two anonymous reviewers for constructive comments. The Fe isotope work was supported by NASA Astrobiology Institute Award NNA04CC04A From Early Biospheric Metabolisms to the Evolution of Complex Systems (to KJE). Support for OJR was provided by a postdoctoral fellowship from the Deep Ocean Exploration Institute at WHOI. This is WHOI contribution number xxxxx.

Supporting Online Material

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Materials and Methods

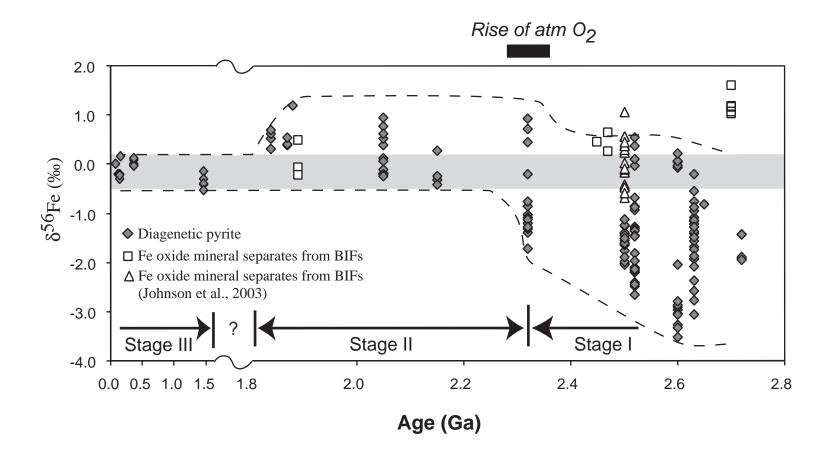
Fig. S1

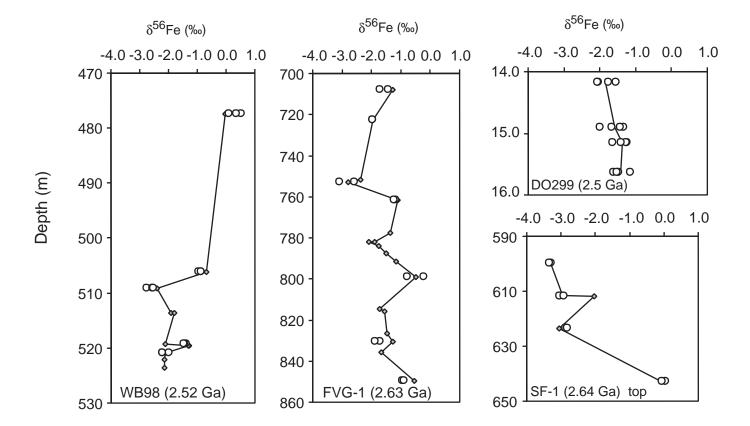
Tables S1, S2, S3

Figure captions

Fig.1: Plot of δ^{56} Fe versus sample age for Fe-sulfides from black shales and Fe-oxides from banded iron formations (BIF). Based on δ^{56} Fe, Fe ocean cycle can be roughly divided into three stages: I) >2.8-2.3 Ga; II) 2.3-1.8 Ga; III) less than 1.7 Ga. Note the scale change between 1.5 to 1.8 Ga. Gray diamonds correspond to Fe isotope composition of pyrite from black shales (Table S2) and open squares and triangles correspond to Fe isotope composition of magnetite and hematite-rich samples from BIFs (Table S3 and Johnson *et al.* (12) respectively). Gray area corresponds to δ^{56} Fe of Fe derived from igneous rocks (at 0.1‰) and hydrothermal sources (ca. -0.5‰) (8). Dashed lines represent the contour lines of maximum and minimum Fe isotope compositions of sedimentary sulfides used to define Stages I to III. The rise of atmospheric oxygen (atm O2) is defined by multiple sulfur isotope analyses of pyrite in the same samples as analyzed for Fe isotopes (3).

Fig. 2: Fe isotope composition of pyrite plotted along stratigraphic sections of black shales. δ^{56} Fe of individual diagenetic pyrite nodules and pyrite grains (open circles) are plotted together with bulk pyrite values (gray diamonds) to illustrate possible Fe isotope heterogeneity within samples. Samples were selected from drill cores WB98 (Gamohaan Formation, Campbellrand Subgroup, Griqualand West Basin, South Africa, ca. 2.52 Ga); FVG-1 (Roy Hill Member of the Jeerinah Formation, upper part of the Fortescue Group, Hamersley Basin, Western Australia, ca. 2.63 Ga); DO299 (Mount McRae Shale, Mount Whaleback Mine, Newman, Hamersley Basin, Western Australia, ca. 2.5 Ga); and SF-1 (Lokammona Formation, Schmidtsdrift Group, Griqualand West Basin, South Africa, ca. 2.64 Ga).





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Supporting Online Material

Materials and Methods

1. Analytical methods

Sample preparation: Clean rock chips were crushed between two plexiglass discs inside a Teflon bag using a hydraulic press. Grains less than 5 mm in size were collected between 500 μm and 1.0 mm sieves and sulfide minerals were isolated by hand-picking under binocular microscope. For each sample, 15 to 20 grains corresponding to weight of 15 to 50 mg were picked in order to obtain a representative sulfide component. To test possible sample heterogeneity, we also separated well developed pyrite nodules, which were individually analyzed. Care was taken to select sulfide grains without organic carbon inclusions but, in some cases, inclusions were unavoidable and their abundance in each sample was qualitatively estimated. For Phanerozoic black shale samples containing only finely disseminated pyrite crystals and framboids, we used chemical leaching procedures to extract pyrite-Fe for Fe isotope analysis. The leaching procedure includes dissolution of silicates with 5 ml of concentrated HF followed by several leaching steps with 50% HCl. The solid residue contains mainly organic carbon and sulfides that are finally dissolved with concentrated HNO₃.

Chemical purification: Samples were weighted in 15 mL Teflon beakers and dissolved using 5 mL of concentrated Trace-metal grade HNO₃. After evaporation on a hot plate at 60°C,

complete dissolution and Fe oxidation was achieved by a second evaporation step using 5mL of *aqua regia*. Dry residue is then dissolved in 5 mL of 6N HCl by heating at 40°C in a closed vessel and ultrasonication. Trace organic carbon occurring mostly as impure inclusions within sulfides was not attacked by this procedure. After centrifugation and separation of C-rich material, a precise solution volume, corresponding to 2500 µg of Fe, was purified on Bio-Rad AG1X8 anion resin (2.5 mL wet bed). After 20 mL of 6N HCl was passed through the column to remove the matrix, 20 mL of 0.12N HCl was used to elute Fe. Eluted solution was then evaporated to dryness and dissolved with 1% HNO₃ to obtain 5 mL of 500 ppm solution.

Mass spectrometry: ⁵⁶Fe/⁵⁴Fe, ⁵⁷Fe/⁵⁴Fe, and ⁵⁷Fe/⁵⁶Fe were determined with a Finnigan *Neptune* multicollector inductively coupled plasma mass spectrometry (MC-ICPMS) operated at Woods Hole Oceanographic Institution (WHOI). The *Neptune* instrument permits high precision measurement of Fe isotope ratios without argon interferences by using high-mass resolution mode (*S1*, *S2*). Mass resolution power of about 8000 (medium resolution mode) was used to resolve isobaric interferences, such as ArO on ⁵⁶Fe, ArOH on ⁵⁷Fe, and ArN on ⁵⁴Fe.

Instrumental mass bias is corrected using Ni isotopes as internal standard. This method, which has been proved to be reliable for the Neptune instrument, involves deriving the instrumental mass bias from simultaneous measuring Ni standard solution (S2). We also used the "sample-standard bracketing" technique to correct for instrumental mass discrimination by normalizing Fe isotope ratios to the average measured composition of the standard that was run before and after the sample (S3-S5). Because the later method may be prone to matrix effects, combination of both methods permits to verify the absence of instrumental artifacts generated by residual matrix elements after chemical purification of samples.

Fe isotope compositions are reported relative the Fe-isotope standard IRMM-14 using the following notations:

$$\delta^{56} Fe = 1000 * [(^{56} Fe/^{54} Fe)_{sample}/(^{56} Fe/^{54} Fe)_{IRMM-14} - 1]$$

$$\delta^{57} Fe = 1000 * [(^{57} Fe/^{54} Fe)_{sample}/(^{57} Fe/^{54} Fe)_{IRMM-14} - 1]$$

$$\delta^{57/56} Fe = 1000 * [(^{57} Fe/^{56} Fe)_{sample}/(^{57} Fe/^{56} Fe)_{IRMM-14} - 1]$$

Sample and standard solutions were diluted to 1.5 ppm of Fe and Ni and introduced into the plasma using a double quartz spray chamber system (cyclonic and double pass) and a microconcentric PFA nebulizer operating at a flow rate of about 200 µl/min. ⁵⁴Fe, ⁵⁶Fe, ⁵⁷Fe, ⁶⁰Ni, and ⁶²Ni isotopes were counted on the Faraday cups using the medium mass resolution mode as described previously (*S1*, *S2*). Baseline corrections were made before acquisition of each data block by completely deflecting the ion beam. Samples were analyzed once or twice and the internal precision of the data are given at 95% confidence levels based on the isotopic deviation of the bracketing standards analyzed during the same analytical session.

Several georeference materials (S6), including one banded iron formation (IF-G), Hawaiian Basalt (BHVO-1) and Cr-rich ultramafic rock (DTS-1) were measured and results are given in Table S1. Based on duplicated chemical purification and isotope analysis, the long term external reproducibility is 0.10% for δ^{56} Fe values, 0.14% for δ^{57} Fe, and 0.07% for $\delta^{56/57}$ Fe (2 standard deviations). Results for mantle-derived rocks give δ^{56} Fe values between 0.07% to 0.11% relative to IRMM-14 that is statistically indistinguishable from previous studies (S4, S5). IF-G gave δ^{56} Fe value of 0.64% similar, within uncertainty, to the 0.67% value obtained by Dauphas et al. (S7). Fe isotope composition of samples are given in Table S2 (Black Shales) and Table S3 (Iron Formations) including numerous duplicate analysis which confirm the overall

analytical precision given above. For all samples, Fe isotope compositions determined using the standard-sample bracketing method and using Ni as internal standard agree well within the uncertainty (Table S1, S2 and S3) confirming further the lack of matrix effects. When Ni was used as internal standard, Cr was not monitored during Fe isotope measurements. However, the analysis of Cr-rich samples (DTS-1) and measurement of Cr concentration similar to the instrumental background for selected samples together with the good relationship between ⁵⁷Fe/⁵⁶Fe *vs.* ⁵⁶Fe/⁵⁴Fe (see below) suggest that Cr is entirely separated from Fe during chemical purification.

The relationships between δ^{56} Fe, δ^{57} Fe and $\delta^{56/57}$ Fe for the samples analyzed in this work are presented in Fig.S1 and all Fe-isotope values plot on a single mass fractionation line. The slope of 0.674 ± 0.007 derived from the relationship between δ^{56} Fe and δ^{57} Fe (Fig. S1) is statistically indistinguishable from that expected for equilibrium (0.678) and kinetic (0.672) fractionations and does not permit to attribute any specific mechanism of isotope fractionation as suggested previously (*S2*, *S8*).

2. Description of the studied material

Pyrite grains in organic-rich shales and Fe-oxides in BIFs have been selected for this study. Pyrite occurs as nodules, disseminated grains, and laminated seams up to several centimeters in thickness. In general, pyrite nodules are 250 µm to several cm in diameter with variable amounts of C-rich inclusions. They either have no internal structure or are composed of concentrically laminated, fine-grained pyrite or bladed pyrite crystals. The outer part of the nodules is commonly overgrown by euhedral pyrite crystals. Early diagenetic origin for most pyrite is supported by the occurrence of laminae in the shale bending around nodules. Pyrite

nodules often display complex features such as multiple-growth bands or composite nodules formed by coalescence of several nodules. Dissolution and reprecipitation of primary sulfide nodules could have happened in some samples and likely resulted in formation of massive pyrite patches, up to several cm in thickness, often characterized by euhedral grains free of C-rich inclusions. Short description of each sulfide sample is given in the Table S2. Geological setting and age constraints are summarized below.

2.1. Proterozoic samples

a) ca. 1.47 Ga Newland Formation, the lower Belt Supergroup, Horse Prairie area, Montana, USA

The minimum age for this formation is constrained by the 1469 ± 2.5 Ma and 1457 ± 2 Ma zircon ages of younger mafic sills (S9). The formation, consisting of intraclast conglomerates, silty black shales with nodular pyrites, dolomitic silts and dolomitic gray shale beds, was deposited in deeper water environment below wave base (S10). The lower Belt Supergroup was deposited during the rifting stage of the Belt basin. Samples in the drill core SC-93 were collected in the lower sulfide zone of the lower part of the Newland Formation, in very thin-bedded pyritic black shale unit composed of few beds of intraformational conglomerate.

b) ca. 1.84 Ga Maraloou Formation, southeastern Capricorn Orogen, Australia

The minimum age of the formation is constrained by the 1843 ± 10 Ma SHRIMP U-Pb monazite age of pepperites in the contact zone with subvolcanic dolerite sills (S11). It belongs to the Mooloogool Subgroup of the Yerrida Group and consists of up to 1 km thick sequence of carbonaceous shale with nodules and bands of pyrite and siltstone with minor sandstone and carbonate deposited in restricted marine to lacustrine settings of the rift basin. Rocks experienced very low metamorphic grade. Samples were collected from the diamond drill core KDD1.

c) ca. 1.88 Ga Dunn Creek Slate, Iron Crystal Falls Range, MI, USA

The Dunn Creek Slate belongs to the Paint Creek River Group which has no direct age constraints. However, the Riverton Iron Formation overlying the Dunn Creek Slate is considered correlative with the Gunflint and Negaunee Iron formations. The latter formations produced similar U-Pb TIMS and SHRIMP zircon ages 1878 ± 2 and 1874 ± 9 Ma, respectively (S12). The unit was deposited in open-marine either retroarc or foredeep Animikie basin. Samples were collected from the drill core DL-92-1 which passed the uppermost part of the Dunn Creek Slate (Wauseca Pyritic Member) consisting of laminated graphitic pyritic slate overlying grey sericitic slate and siltstone (S13).

d) ca. 1.82 Ga Virginia Formation, Animikie Group, Animikie Basin, Minnesota, USA

The Virginia Formation belongs to the Animikie Group, gradationally overlies the Biwabik Iron Formation and was deposited in either retro-arc or fore-deep Animikie Basin. The age of this unit is constrained by the 1821 ± 16 Ma U-Pb SHRIMP zircon age of the ash-bed in the lower part of the correlative Rove Formation in the Gunflint Range that overlies the 1878 ± 2 Ma Gunflint Iron Formation (S14, S15). The Virginia Formation consists of upward-coarsening sequence of laminated mudstone, siltstone, and fine-grained greywacke (S16). Samples of carbonaceous mudstones with pyrite nodules were collected from the recently drilled drill hole VHD-00-1 in the Virginia Horn area which experienced very low metamorphic grade.

e) ca. 2.1-2.0 Ga Francevillian Series, Gabon

The age of the Francevillian Series is poorly constrained by 2036 ± 79 Ma and 2099 ± 115 Ma Sm-Nd ages of clay minerals (S17) and the 1950 ± 40 Ma age of the Oklo natural fission reactor (S18). Carbonates in the lower part of the Francevillian Series have δ^{13} C values ranging from +2.1 to +6.3 ‰ (S19) suggesting deposition during the final stage of the ca. 2.22-2.06 Ga

carbon isotope excursion (S20). The end of this carbon isotope excursion is bracketed on the Fennoscandian Shield and elsewhere between 2.11 and 2.06 Ga. The lower FB member consists of black shales with pyrite and was deposited in an open-marine setting. The succession experienced only low metamorphic grade. Samples were collected from the drill core OKP that was collared on the Okouma plateau.

f) ca. 2.1-2.0 Ga upper Zaonezhskaya Formation, lower Ludikovian Series, Karelia, Russia

The age of the Zaonezhskaya Suite is constrained by its unconformable stratigraphic position in the Lake Onega area above carbonates of the upper Jatulian Series that carry high 13 C-enrichment typical for the ca. 2.2-2.1 Ga carbon isotope excursion (S20). The minimum age is constrained by 1980 ± 27 Ma Sm-Nd mineral isochron age of gabbro intrusion that cuts the overlying unit (S21). The formation experienced greenschist facies metamorphism (S22). The Zaonezhskaya Formation consists of a 1500-m-thick sequence of basaltic tuffs, organic-rich siltstones and mudstones, and cherts with subordinate dolostones. Samples were collected from the drill cores 19, 159, and 5190 from various depths.

g) ca. 2.2-2.1 Ga Sengoma Argillite Formation of the Pretoria Series, Lobatse, Botswana

The Sengoma Argillite Formation has no direct geochronologic constraints, however it is likely correlative with the Silverton Formation of the Pretoria Group in South Africa (S23). The Silverton Formation is intruded by the 2061 \pm 2 Ma Bushveld Complex (S24). The maximum age is constrained by the underlying 2.22 Ga Hekpoort-Ongeluk volcanics, and 2316 \pm 7 Ma Re-Os isochron pyrite age for the Rooihoogte-lower Timeball Hill formations (S25). Further constrains are provided by the overlying carbonates that are 13 C-enriched. We therefore infer that the Sengoma Argillite Formation was deposited during the 2.2-2.1 Ga carbon isotope excursion. The Sengoma Argillite Formation is up to 700 m in thickness upward-shallowing sequence and consists of carbonaceous argillite with pyrite, chert, siltstone, and fine-grained quartzite.

Argillites often have convoluted bedding and sandstone dikes. It was deposited in deep openmarine environment and experienced greenschist facies metamorphism. Samples were collected from the drill core STRAT 2.

h) 2.32 Ga Rooihoogte and Timeball Hill formations, Lower Pretoria Group, Transvaal Basin, South Africa

These units were deposited in a deltaic part of the open-marine Transvaal Basin between the second and third glacial events of the early Paleoproterozoic (S26). Their age is well-constrained by the 2316 ± 7 Ma Re-Os isochron pyrite age (S25). The formations experienced lower greenschist facies metamorphism and consist of four upward-shallowing cycles of black shale with pyrite, siltstone, and sandstone. The most organic- and pyrite-rich parts of the sequence are in the upper part of the Rooihoogte Formation and the lower part of the Timeball Hill Formation. For description of studied pyrite see Bekker et al. (S27). Samples were collected from the drill cores of EBA-series that were collared in the Potchefstroom area, western part of the Transvaal Basin and from the drill core PA-13 collared in the Penge area, eastern part of the Transvaal Basin.

2.2. Archean Samples

a) ca. 2.5 Ga Mount McRae Shale, Mount Whaleback Mine, Newman, Hamersley Basin, Western Australia

The age of the Mount McRae Shale is bracketed by the U-Pb SHRIMP zircon age of 2561 \pm 8 Ma for the Crystal Rich Tuff of the Wittenoom Dolomite and by the 2479 \pm 3 Ma age for the S9 macroband of the Dales Gorge Member of the Brockman Iron Formation (S28, S29). The Mount McRae Shale was deposited in an open-marine setting and consists of carbonaceous pyritic shale and minor chert layers and nodules (S30). Based on the presence of graded beds of

massive and plane-laminated, quartz-sericitic sandstones with tabular shale clasts and erosional bases, dolomitized limestones with floatstones and wackestones in this unit, it was interpreted as deposited by turbidity currents (*S30*). Samples were collected from the drill hole DDH44 collared near Paraburdoo (Colonial Chert Member; upper part of the Mount McRae Shale; AMG zone 50, 0560534mE, 7429955mN) and the drill hole DO299 collared in the Whaleback Mine, Newman, Western Australia (AMG zone 50, 0722600mE, 7412900mN).

b) ca. 2.52 Ga Gamohaan Formation, Campbellrand Subgroup, Griqualand West Basin, South Africa

The Gamahaan Formation is the uppermost unit of the Campbellrand Subgroup immediately underlying the ca. 2465 Ma Kuruman Iron Formation. The age of the Gamohaan Formation is well-constrained by the TIMS U-Pb zircon age for the ash bed within this unit at 2521 ± 3 Ma (S31). The formation was deposited on the open continental margin of the Kaapvaal Craton and consists of peritidal, subtidal, and basinal siliciclastic and carbonate rocks with various microbialite assemblages and abundant inorganic precipitates, carbonaceous shale with pyrite nodules, tuff, chert, and chert and dolostone breccia (S32). Carbonaceous shale is particularly abundant near the top of the formation and contains uncompacted laminated mats and it is interpreted to be formed in deep subtidal environments (S32). Samples were collected from the drill hole WB-98 from the Kuruman area.

c) ca. 2.6 Ga Carawine Dolomite, Hamersley Group, Hamersley Basin, Western Australia

The Carawine Dolomite is present along the northeastern edge of the Pilbara Craton in the area separated from the main outcrop of the Hamersley Group. The age of the Carawine Dolomite is not constrained directly by published geochronologic data, however ash bed at the base of the Carawine Dolomite has SHRIMP U-Pb zircon age of 2.63 Ga (Bryan Krapez, pers. com., 2004). The maximum age of the Carawine Dolomite is constrained by the SHRIMP U-Pb

zircon age of 2764 ± 8 Ma for the Koongaling Volcanics at the base of the underlying Fortescue Group in this area (S33). It overlies the Lewin Shale which is considered correlative with the ca. 2.65 Ga Jeerinah Formation. The Carawine Dolomite contains large stromatolitic domes, laminites and clastic carbonate strata, oncolites, wave ripples, roll-up structures, and locally preserved evaporite crystal pseudomorphs, oölites, and pisolites indicating deposition in shallowwater platform environment (S34). The Carawine Dolomite consists almost entirely of dolomite with thin argillite partings and beds and chert nodules (Simonson et al., 1993). Samples of the Carawine Dolomite were collected from the drill core RHDH2A.

d) ca. 2.65 Ga Lewin Shale / Jeerinah Formation, upper part of the Fortescue Group, Hamersley Basin, Western Australia

The age of the Jeerinah Formation and correlative Lewin Shale in the eastern part of the Pilbara Craton is bracketed by the 2629 ± 5 Ma SHRIMP U-Pb zircon age of andesitic ignimbrite at the top of the Roy Hill Shale Member, the upper Jeerinah Formation (S28) and 2715 ± 2 Ma and 2713 ± 3 Ma SHRIMP U-Pb zircon ages for felsic tuffs within the underlying Maddina Basalt (S35). The ash bed with the similar age is also present at the base of the Carawine Dolomite (B. Krapez pers. com., 2004). Arndt et al. (S33) obtained 2684 ± 6 Ma and 2690 ± 16 Ma SHRIMP U-Pb zircon ages for the andesitic ignimbrite and tuffaceous sandstone of the upper Jeerinah Formation. The Jeerinah Formation was deposited during the transition from the rift to drift stage in an open-marine setting and consists of upward-fining sequence of sandstone, siltstone, shale, and carbonaceous pyritic shale with minor chert, dolomite, and mafic and felsic volcanic and volcanoclastic rocks (S36). Samples of the Lewin Shale were collected from the drill core RHDH2A whereas the Roy Hill Member samples were collected from the drill core FVG-1.

e) ca. 2.6 Ga Lokammona Formation, Schmidtsdrif Subgroup, Griqualand West Basin, South Africa

The age of the Schmidtsdrif Subgroup of the Ghaap Group, the Griqualand West Basin is only constrained by the Pb-Pb zircon age of 2642 ± 3 Ma for the Vryburg Formation, which is at the base of the subgroup (Walvaren et al., unpubl. in Nelson et al., (S28)). The unconformably underlying Ventersdorp Supergroup provides the maximum age constraint of 2709 ± 4 Ma for the Schmidtsdrif Subgroup SHRIMP U-Pb zircon age for the Makwassie Quartz Porphyry Formation (S37)). The ash bed in the lower part of the conformably overlying Campbellrand Subgroup provides minimum age constraint for the Schmidtsdrif Subgroup of 2588 ± 6 Ma (SHRIMP U-Pb zircon age; (S38)). The Lokammona Formation forms the uppermost part of the subgroup and was deposited in deep-water environment on the open-marine margin and consists of upward-fining cycles of siltstone and carbonaceous argillite with minor amounts of chert and carbonate grainstones (S39). Carbonaceous argillites contain nodular, disseminated, and partially recrystallized pyrite. Samples were collected from the drill hole SF-1 collared in the Vryburg area.

f) ca. 2.72 Ga Deer Lake greenstone sequence, MN, USA

The Deer Lake greenstone sequence is located in the north-central Minnesota and belongs to the western Wawa Subprovince of the Superior Province. No direct age constraints are available for this Archean succession, although it is considered correlative with the Soudan Belt of the Vermillion District and its extension to Canada, the Shebandowan greenstone Belt, both of which developed between 2.75 and 2.67 Ga (S40, S41). The sequence experienced metamorphism of greenschist facies and consists of interlayered mafic to intermediate volcanics, greywackes, and carbonaceous slates and argillites with pyrite nodules up to 2 cm in diameter (S42). Samples were collected from the drill holes 26503 and 26516.

g) ca. 2.72 Ga Helen Iron Formation, Michipicoten Group, MacLeod and Sir James Mines, Wawa, Ontario, Canada

The age of the Helen Iron Formation that includes sideritic stromatolites and organic-rich shales with pyrite nodules up to 5 cm in diameter is well-constrained by the TIMS U-Pb zircon age of felsic tuffs immediately underlying the Iron Formation at 2749 ± 2 Ma at Helen Iron Range in Wawa (S43) whereas in Goudreau intermediate to felsic tuffs below the correlative Morrison No. 1 Iron Formation have TIMS U-Pb zircon age 2729 ± 2 Ma (S44). The overlying volcanics have TIMS U-Pb zircon age of 2696 ± 2 Ma (S43). Deposition of the Helen Iron Formation took place during the hiatus between the second and third cycles of bimodal basalt-rhyolite volcanism at 2.75 and 2.70 Ga, respectively, in island arc to mature arc setting built on continental crust (S45). Small bioherms with complex conical stromatolites occur at the top of the siderite ore body in the Helen Iron Formation at the MacLeod Mine (S46). Secondary pyrite and magnetite develop along the lamina. Carbonaceous argillite with pyrite nodules occur at the top of the Iron Formation (S47). Pyrite nodules have radial and concentric morphology. Stromatolitic siderite was collected in the MacLeod Mine and carbonaceous argillites with pyrite nodules were collected on the north side of the Sir James (=Eleanor) Mine, Eleanor Iron Range.

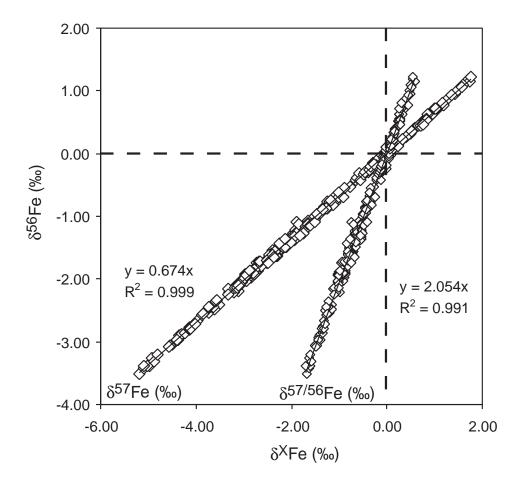


Fig.S1: Three-isotope plot of Fe isotope values of black shale sulfides. Squares show δ^{56} Fe νs . δ^{57} Fe data and diamonds represent δ^{56} Fe νs . $\delta^{57/56}$ Fe data.

Table S1: Duplicated Fe-isotope composition of georeference Materials

Name	Method	δ^{56} Fe	2σ	δ^{57} Fe	2σ	δ ^{57/56} Fe	2-		
							2σ		
BHVO-1 #1	(a)	0.08	0.13	0.10	0.22	0.01	0.10		
BHVO-1 #2	(a)	0.05	0.12	0.11	0.19	0.04	0.05		
duplicate	(b)	0.09	0.14	0.16	0.20	0.06	0.07		
BHVO-1 #3	(a)	0.18	0.13	0.26	0.10	0.10	0.16		
duplicate	(b)	0.17	0.19	0.22	0.20	80.0	0.10		
BHVO-1 #4	(b)	0.19	0.07	0.22	0.19	0.02	0.04		
duplicate BHVO-1 #5	(b)	0.16	0.13	0.19	0.19	0.05	0.07		
	(b)	0.02	0.15	0.04	0.24	0.03	0.10		
duplicate	(b)	0.07	0.06	0.07	0.08	0.00	0.04		
BHVO-1 #6	(b)	0.05	0.16	0.14	0.13	0.06	0.09		
BHVO-1 #7	(a)	0.06	0.16	80.0	0.16	0.03	0.06		
duplicate BHVO-1 #8	(b)	0.12	0.10	0.22	0.20	0.10	0.12		
	(a)	0.11	0.04	0.13	0.14	0.01	0.07		
duplicate	(a)	0.02	0.06	0.06	0.14	0.02	0.08		
BHVO-1 #9 BHVO-1 #10	(b)	0.03	0.16	0.08	0.13	0.03	0.09		
BHVO-1 #10	(b)	0.04	0.16	0.06	0.13	0.03	0.09 0.06		
	(a)	0.15	0.09	0.17	0.03	0.04	0.06		
duplicate BHVO-1 #12	(b)	0.16 0.07	0.14 0.15	0.18 0.17	0.20 0.17	0.06 0.10	0.07		
duplicate	(a) (b)	0.07	0.13	0.17	0.17	0.10	0.03		
BHVO-1 #13	(a)	0.09	0.07	0.21	0.10	0.12	0.04		
duplicate	(a) (b)	0.16	0.13	0.18	0.20	0.11	0.17		
BHVO-1 #14	(a)	0.14	0.00	0.05	0.12	0.00	0.03		
BHVO-1 #15	(a) (a)	0.14	0.12	0.13	0.27	0.01	0.13		
duplicate	(b)	0.13	0.00	0.20	0.20	0.02	0.21		
BHVO-1 #16	(b)	0.14	0.11	0.16	0.14	0.03	0.03		
duplicate	(a)	0.14	0.11	0.15	0.16	0.02	0.10		
duplicate	(b)	0.12	0.11	0.20	0.20	0.09	0.09		
BHVO-1 #17	(a)	0.12	0.08	0.17	0.02	0.03	0.03		
duplicate	(b)	0.21	0.08	0.29	0.13	0.10	0.08		
BHVO-1 #18	(a)	0.07	0.13	0.08	0.40	0.01	0.25		
duplicate	(b)	0.03	0.08	0.00	0.13	-0.03	0.08		
BHVO-1 #19	(a)	0.14	0.22	0.17	0.00	0.04	0.22		
duplicate	(b)	0.20	0.11	0.23	0.14	0.05	0.09		
BHVO-1 #20	(a)	0.06	0.11	0.08	0.18	0.02	0.13		
duplicate	(b)	0.11	0.11	0.15	0.14	0.03	0.09		
BHVO-1 All data	(3)	0.11	0.11	0.15	0.13	0.05	0.07		
DTS-1 #1	(a)	0.11	0.08	0.17	0.11	0.10	0.03		
duplicate	(b)	0.03	0.14	0.04	0.20	0.06	0.07		
DTS-1 #2	(b)	0.02	0.16	0.04	0.13	0.02	0.09		
DTS-1 All data	(/	0.06	0.10	0.08	0.15	0.06	0.09		
IF-G#1	(b)	0.68	0.06	1.02	0.08	0.35	0.04		
duplicate	(b)	0.62	0.16	0.95	0.13	0.33	0.09		
IF-G#2	(a)	0.63	0.07	0.94	0.07	0.30	0.04		
duplicate	(b)	0.63	0.08	0.92	0.12	0.26	0.05		
IF-G All data	(- /	0.64	0.06	0.96	0.09	0.31	0.08		
(a): Instrumental mass higs corrected using the "standard-sample bracketing"									

(a): Instrumental mass bias corrected using the "standard-sample bracketing" method; (b): Instrumental mass bias corrected using Nickel standard solution as an internal standard.

Table S2: Fe-isotope composition of pyrite from Black Shales									
Sample	Met	δ^{56} Fe	2σ	δ^{57} Fe	2σ	$\delta^{57/56}$	2σ	org-C	Description
	hod					Fe			
Kentucky Black Sh		-					-		
Clay C. 510-519 #1	(a)	0.01	0.19	-0.02		-0.02		+++	500μm framboids Py
Clay C. 510-519 #2	(a)	-0.04	0.07	-0.06		-0.04		+	Massive Py (in veins)
duplicate	` '	-0.01	0.06	0.01		-0.01	0.04		
duplicate	٠,,	-0.04	0.08	0.01	0.10	0.03	0.05		
duplicate	(b)	-0.02	0.16	0.06	0.13	0.04	0.09		
Clay C. 193-200	(a)	-0.01	0.09	-0.08		-0.03	0.13		Bulk Black Shale (Leach)
Clay C. 232-238	(b)	0.09	0.08	0.08	0.12	0.03	0.05		Bulk Black Shale (Leach)
2000NAS	(b)	-0.02	0.13	0.02	0.19	0.02	0.07		Bulk Black Shale (Leach)
Black Shales, Egyp	ot (De	evonian)						
B4,202	(b)	0.12	0.13	0.20		0.09	0.07		Bulk Black Shale (Leach)
Black Shales , Egy	pt (E	arly Cre	taceo	us, Aptı	ian)				
KB40	(b)	-0.20	0.13	-0.32	0.19	-0.12	0.07		Bulk Black Shale (Leach)
Black Shales , Egy	pt (La	ate Cret	taceou	is, Cam	paniai	n)			
Abu-2	(b)	0.01	0.08	0.01	0.12	-0.02	0.05		Bulk Black Shale (Leach)
Black Shales , Egy	pt (Ji	urassic))						
B2-4	(a)	0.16	0.14	0.23	0.21	0.05	0.06		Bulk Black Shale (Leach)
Kimmeridge Bay B	lack	Shales	(Late .	Jurassi	c); Ma	thews	et al.,	2004	
BLN-A		-0.21	•	-0.31					Nod. Py
BLN-C		-0.30		-0.45					Nod. Py
CLN-1		-0.23		-0.34					Nod. Py
									•
ca. 1.47 Ga Newlan	d Fo	rmation	, lowe	r Belt S	uperg	roup,	Horse	Prairi	ie area, Montana, USA
SC-93, 1795	(a)	-0.14	0.10	-0.30	0.14	-0.04	0.03	+	Massive & Fine-g Py
duplicate	(b)	-0.25	0.15	-0.37	0.24	-0.11	0.10		
SC-93, 1802 a	(a)	-0.31	0.09	-0.48	0.16	-0.14	0.13	+	Massive & Fine-g Py
SC-93, 1802 b	(b)	-0.42	0.08	-0.65	0.10	-0.21	0.04	+	Massive & Fine-g Py
SC-93, 1809 a	(b)	-0.39	0.14	-0.62	0.20	-0.21	0.07	+	Massive & Fine-g Py
SC-93, 1809 b	(b)	-0.53	0.06	-0.78	0.08	-0.27	0.04	+	Massive & Fine-g Py
duplicate	. ,	-0.64	0.04	-0.98	0.06	-0.34	0.04		3 ,
ca. 1.84 Ga Maralo	. ,		n, sou	theaste	rn Ca	oricorn	Orog	jen, A	ustralia
KDD1(1), 98.75	(a)	0.31	0.09	0.48	-	0.17	_	O	Massive Py along layers
duplicate	٠,,	0.34	0.14	0.49	0.20	0.15	0.07		, , ,
duplicate		0.24	0.08	0.33	0.10	0.13	0.04		
KDD1(3), 148.9	(a)	0.52	0.10	0.77	0.14	0.28	0.03	0	Massive Py along layers
duplicate		0.50	0.08	0.73	0.08	0.25	0.04		massive y among region
KDD1(6), 297.15	(b)	0.62	0.09	0.88	0.19	0.26	0.04	++	Fine-g Py in Nodules
KDD1, 273.7	(a)	0.69	0.08	1.05	0.10	0.35	0.08	+++	Massive Py in Nodules
duplicate	٠,,	0.62	0.09	0.92	0.19	0.29	0.04		accirc i y i caa.cc
ca. 1.82 Ga Virginia								in. Min	nesota. USA
VHD-001, 1625 a	(a)	0.39	0.09	0.57	0.07	0.15	0.05	0	Massive & Fine-g Py
duplicate	٠,,	0.35	0.05	0.53	0.24	0.16	0.10	J	assive a rine gry
VHD-001, 1625 b	(b)	0.33	0.13	0.60	0.24	0.10	0.10	0	Massive & Fine-g Py
VHD-001, 1023 b	(b)	0.42	0.14	0.81	0.20	0.19	0.07		Massive Ry along layers
	. ,	0.62	0.14	0.94	0.20	0.20		0	wassive ry along layers
duplicate	(n)	0.02	0.05	0.34	0.00	U.3Z	0.04		

```
ca. 1.88 Ga Dunn Creek Slate, Iron Crystal Falls Range, MI, USA
DL-92-1, 550
                    (a)
                          1.19
                                0.26
                                       1.72
                                              0.33 0.55
                                                          0.07
                                                                      Euh Py
                                0.14
                                       1.74
                                              0.20 0.59
          duplicate (b)
                          1.16
                                                          0.07
          duplicate (a)
                         1.10
                                0.06
                                       1.58
                                              0.09
                                                    0.50
                                                          0.07
          duplicate (b)
                                       1.75
                          1.22
                                0.05
                                              80.0
                                                    0.55
                                                          0.04
ca. 2.1-2.0 Ga Francevillian Series. Gabon
OKP, 67.6 a
                    (b)
                          0.39
                                0.16
                                       0.56
                                              0.19 0.18 0.09
                                                                 +++ Fine-g Py along layers
OKP, 67.6 b
                    (b)
                          0.52
                                0.16
                                       0.76
                                              0.13
                                                    0.24
                                                          0.09
                                                                 +++ Fine-g Py along layers
          duplicate (b)
                         0.51
                                80.0
                                       0.77
                                              0.12 0.25
                                                          0.05
OKP, 69.98
                         0.77
                                0.24
                                       1.19
                                              0.37
                                                    0.45
                                                          0.17
                                                                 +++ Fine-g Py along layers
                    (a)
                                       1.03
          duplicate (a)
                         0.72
                                0.14
                                              0.30 0.31
                                                          0.12
          duplicate (b)
                          0.88
                                0.08
                                       1.27
                                              0.12 0.40 0.05
                                       0.23
OKP, 70.45
                    (b)
                          0.12
                                0.16
                                              0.19
                                                    0.04 0.09
                                                                      Fine-g Py along layers
OKP, 70.6
                                       -0.25
                    (a)
                         -0.19
                                0.08
                                              0.11 -0.09 0.03
                                                                      Fine-g & Massive Py
                                                                      along layers
          duplicate (b)
                         -0.15
                                0.14
                                       -0.21
                                              0.20 -0.07 0.07
          duplicate (b)
                         -0.05
                                0.06
                                       -0.15
                                              0.09 -0.06 0.05
ca. 2.1-2.0 Ga upper Zaonezhskaya Formation, lower Ludikovian Series, Karelia, Russia
159/81.5 a
                                0.10
                                       0.85
                                              0.12 0.27 0.02
                                                                      Euh Pv
                    (a)
                         0.62
                          0.72
                                0.06
                                       0.99
                                              0.08
          duplicate (b)
                                                    0.30
                                                          0.04
                                       -0.09
159/81.5 b
                         -0.11
                                0.08
                                              0.10 -0.02 0.03
                    (a)
                                                                      Fine-g Py
          duplicate (b)
                         -0.06
                                0.06
                                       -0.07
                                              0.08 -0.01 0.04
19/1.62
                         0.06
                                0.06
                                       0.09
                                              0.08 0.02
                                                          0.06
                                                                      Euh Py
                    (a)
          duplicate (b)
                         0.03
                                0.06
                                       0.02
                                              0.08 -0.03 0.04
                                       -0.32
                                              0.08 -0.09 0.06
19/159.5
                    (a)
                         -0.20
                                0.06
                                                                      Euh Py
                         -0.12
                                0.06
                                       -0.22
                                              0.08 -0.08 0.04
          duplicate (b)
5190/158
                         0.94
                                0.06
                                       1.43
                                              0.08 0.51
                                                                      Massive Py
                    (b)
                                                          0.04
                                0.16
                                       1.47
          duplicate (b)
                          1.01
                                              0.13 0.48
                                                          0.09
5190/85
                         -0.24
                                0.05
                                      -0.26
                                              0.08 -0.02 0.03
                    (a)
                                                                      Massive Py
          duplicate (b)
                         -0.29
                                0.06
                                      -0.39
                                              0.08 -0.11 0.04
ca. 2.2-2.1 Ga Sengoma Argillite Formation of the Pretoria Series, Lobatse, Botswana
Strata2, 138
                         -0.32
                                0.14
                                       -0.58
                                              0.20 -0.14 0.07
                                                                      Massive Py in nodules
                    (b)
                                              0.09 -0.15 0.05
                                0.05
                                       -0.43
          duplicate (b)
                         -0.30
Strata2, 192.2
                         -0.25
                                0.14
                                       -0.34
                                              0.22 -0.08 0.09
                    (a)
                                                                  O
                                                                      Euh. Py
                         -0.42
                                0.13
                                       -0.55
                                              0.22 -0.13 0.09
Strata2, 192.8
                    (a)
                                                                      Massive Py in nodules
                                0.05
                                       -0.49
          duplicate (a)
                         -0.33
                                              0.08 -0.17 0.02
          duplicate (b)
                         -0.30
                                0.05
                                       -0.46
                                              0.09 -0.16 0.05
Strata2, 195.0
                         0.27
                                0.05
                                       0.39
                                              0.08 0.10 0.03
                                                                      Euh. Py
                    (a)
          duplicate (b)
                         0.28
                                0.06
                                       0.43
                                              0.08 0.14 0.04
2.32 Ga Rooihoogte and Timeball Hill formations, Lower Pretoria Group, Transvaal Basin,
South Africa
 Timeball Hill Fm.
EBA-1, 1052.0
                    (a)
                         0.45
                                0.05
                                       0.67
                                              0.12 0.24
                                                          0.09
                                                                      Fine-g Py
                                0.07
                                       0.82
                                                    0.28
          duplicate (b)
                         0.56
                                              0.10
                                                          0.04
                         0.93
                                0.05
                                       1.36
                                              0.13 0.44
EBA-1, 1057.93
                    (a)
                                                          0.08
                                                                      Massive Py
          duplicate (b)
                          1.05
                                0.05
                                       1.56
                                              0.09 0.52
                                                          0.05
EBA-1, 1160.2
                         -1.20
                                0.05
                                       -1.81
                                              0.09 -0.62 0.04
                                                                      Massive Py
                    (a)
                                       -2.06
                                              0.20 -0.70 0.07
                         -1.37
                                0.14
          duplicate (b)
          duplicate (a)
                         -1.25
                                0.05
                                      -1.86
                                              0.13 -0.63 0.08
          duplicate (b)
                         -1.15
                                0.05
                                       -1.69
                                              0.09 -0.56 0.05
EBA-3, 1164.8
                         0.72
                                0.05
                                       1.03
                                              0.08 0.25
                    (a)
                                                          0.03
                                                                      Massive & Fine-g Py
                                0.07
                                       1.17
                                              0.10 0.31 0.04
          duplicate (b)
                         0.81
```

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EBA-2/30
                          -1.73
                                 0.05
                                        -2.56
                                               0.08 -0.84 0.03
                                                                        Massive Py
                     (a)
                                        -2.41
           duplicate (b)
                          -1.67
                                 0.16
                                               0.13 -0.73
                                                            0.09
EBA-2/31
                                 0.05
                                        -1.54
                                               0.08 -0.51
                          -1.02
                                                            0.03
                                                                        Fine-g Py
                     (a)
           duplicate (b)
                          -0.96
                                 0.06
                                        -1.47
                                               0.08 -0.51
                                                            0.04
                                        -1.32
EBA-2/32
                     (a)
                          -0.86
                                 0.05
                                               80.0
                                                     -0.45
                                                            0.02
                                                                        Fine-g Py
                          -0.88
                                 0.06
                                        -1.32
                                               0.08 -0.45
           duplicate (b)
                                                            0.04
EBA-2/63
                          -1.40
                                 0.05
                                        -2.11
                                               0.08 -0.70
                                                                        Massive Py
                     (a)
                                                            0.02
           duplicate (b)
                          -1.37
                                 0.06
                                        -2.04
                                               80.0
                                                    -0.68
                                                            0.04
 Rooihoogte Fm.
EBA-3, 1297.5
                                 0.05
                                        -1.89
                                               0.12 -0.54 0.09
                                                                        Massive Py
                          -1.33
                     (a)
           duplicate (b)
                          -1.36
                                 0.07
                                        -1.95
                                               0.10 -0.57
                                                            0.04
EBA-2/67
                     (a)
                          -1.08
                                 0.05
                                        -1.60
                                               80.0
                                                    -0.53
                                                            0.02
                                                                        Massive Py
                                                                    0
                                        -1.51
           duplicate (b)
                          -1.05
                                 0.06
                                               0.08 -0.47
                                                            0.04
                                 0.08
                                        -1.24
EBA-2/55-2
                          -0.76
                                               0.10 -0.44
                                                            0.02
                                                                        Massive Py
                     (a)
                                                                    0
                          -0.68
                                 0.06
                                        -1.08
                                               80.0
                                                    -0.37
           duplicate (b)
                                                            0.04
EBA-2/55-3
                                        -1.72
                                               0.28 -0.54
                     (a)
                          -1.12
                                 0.13
                                                                        Massive Pv
                                        -1.88
           duplicate (b)
                          -1.25
                                 0.08
                                               0.12 -0.59
                                                            0.05
EBA-2/59
                                        -0.29
                          -0.21
                                 0.13
                                               0.18 -0.07
                                                            0.02
                                                                        Massive Py
                     (a)
                          -0.26
                                 0.08
                                        -0.37
                                               0.12 -0.12 0.05
           duplicate (b)
EBA-2/60
                          -1.28
                                        -1.97
                                               0.18 -0.68
                     (a)
                                 0.13
                                                            0.02
                                                                        Massive Py
           duplicate (b)
                          -1.23
                                 0.08
                                        -1.91
                                               0.12 -0.68 0.05
ca. 2.5 Ga Mount McRae Shale, Mount Whaleback Mine, Newman, Hamersley Basin, Western
Australia
DO299, 14.2 #1
                     (a)
                          -2.03
                                 0.07
                                        -3.01
                                               0.20 -0.99
                                                            0.12
                                                                        Nod py (2mm)
                          -2.06
                                 0.04
                                        -3.07
                                               0.06 -1.03 0.04
           duplicate (b)
DO299, 14.2 #2
                     (a)
                          -2.05
                                 0.04
                                        -3.07
                                               0.08 -1.00
                                                            0.04
                                                                        Nod py (4mm)
                          -2.00
                                 0.04
                                        -2.97
                                               0.06
                                                    -0.96
           duplicate (b)
                                                            0.04
                                        -2.72
DO299, 14.2 #3
                          -1.75
                                 0.04
                                               0.09 -0.96
                                                            0.04
                                                                        Nod py (4mm)
                     (a)
                          -1.73
                                 0.04
                                        -2.67
                                               0.06 -0.94
           duplicate (b)
                                                            0.04
                                                                        Nod py (4mm)
DO299, 14.2 #4
                     (a)
                          -1.54
                                 0.28
                                        -2.39
                                               0.36 -0.80
                                                            80.0
                                        -2.24
                          -1.55
                                 0.05
                                               0.08 -0.69
           duplicate (a)
                                                            0.04
                                        -2.29
                          -1.58
                                 0.08
                                               0.12 -0.73
           duplicate (b)
                                                            0.05
DO299, 14.95 #1
                          -1.41
                                 0.05
                                        -2.10
                                               0.05 -0.70
                                                            0.03
                                                                        Nod py (2mm)
                     (a)
           duplicate (b)
                          -1.37
                                 0.05
                                        -2.08
                                               0.08 -0.72
                                                            0.05
                          -1.65
                                        -2.38
DO299, 14.95 #2
                                 0.05
                                               0.06
                                                    -0.71
                                                                        Nod py (4mm)
                     (a)
                                                            0.03
           duplicate (b)
                          -1.60
                                 0.05
                                        -2.34
                                               0.08 -0.73
                                                            0.05
DO299, 14.95 #3
                                               0.06 -0.91
                                 0.05
                                        -2.89
                                                                        Nod py (4mm)
                     (a)
                          -1.98
                                                            0.03
                                        -2.79
                                                    -0.91
           duplicate (b)
                          -1.89
                                 0.05
                                               80.0
                                                            0.05
                                        -1.76
DO299, 15.2 #1
                     (a)
                          -1.25
                                 0.05
                                               0.08 -0.52 0.06
                                                                        Nod py (3mm)
                                        -1.78
           duplicate (b)
                          -1.25
                                 0.05
                                               0.08 -0.52
                                                            0.05
                                        -2.01
DO299, 15.2 #2
                     (a)
                          -1.39
                                 0.05
                                               0.08 -0.63
                                                            0.06
                                                                        Nod py (3mm)
                                        -2.01
           duplicate (b)
                          -1.38
                                 0.05
                                               0.08 -0.63
                                                            0.05
                                        -2.30
                                                    -0.70
DO299, 15.2 #3
                     (a)
                          -1.63
                                 0.06
                                               0.09
                                                            0.09
                                                                        Nod py (3mm)
                          -1.62
                                 0.05
                                        -2.33
                                                    -0.72
           duplicate (b)
                                               80.0
                                                            0.05
DO299, 15.2 #4
                     (a)
                          -1.23
                                 0.25
                                        -1.78
                                               0.17 -0.62
                                                            0.05
                                                                        Nod py (3mm)
                          -1.15
                                 0.07
                                        -1.70
                                               0.10 -0.62
           duplicate (b)
                                                            0.04
DO299, 15.7 #1
                     (a)
                          -1.46
                                 0.04
                                        -2.09
                                               0.01
                                                    -0.68
                                                            0.09
                                                                        Nod py (2mm)
           duplicate (b)
                          -1.48
                                 0.05
                                        -2.16
                                               0.08 -0.71
                                                            0.05
                          -1.60
                                 0.08
                                        -2.41
                                                     -0.81
                                                                        Nod py (2mm)
DO299, 15.7 #2
                     (a)
                                               0.11
                                                            0.10
                                        -2.28
                                               0.08 -0.78
           duplicate (b)
                          -1.50
                                 0.05
                                                            0.05
                                        -2.29
                                               0.11 -0.78
DO299, 15.7 #3
                     (a)
                          -1.51
                                 0.05
                                                            0.10
                                                                        Nod py (2mm)
                                        -2.25
           duplicate (b)
                          -1.47
                                 0.05
                                               80.0
                                                    -0.79 0.05
```

```
DO299, 15.7 #4
                    (a) -1.13
                                0.03 -1.73 0.12 -0.56 0.10
                                                                      Nod py (2mm)
          duplicate (b) -1.06
                                0.19
                                      -1.66
                                              0.19 -0.59 0.10
ca. 2.5 Ga Colonial Chert Member, Mount McRae Shale, Hamersley Basin, Western Australia
DDH44, 516.44
                    (a)
                         -1.89
                                0.11
                                       -2.89
                                              0.03 -1.00 0.05
                                                                      Massive Pv
                                       -2.76
          duplicate (a)
                         -1.91
                                0.08
                                              0.15 -0.87 0.13
          duplicate (b)
                         -1.87
                                0.05
                                      -2.71
                                              0.09 -0.86 0.05
ca. 2.52 Ga Gamohaan Formation, Campbellrand Subgroup, Griqualand West Basin, South
Africa
WB98, 477.50
                    (a)
                         -0.03
                                0.08
                                       0.06
                                              0.11
                                                    0.05 0.12
                                                                      Massive Py in Nodules
          duplicate (b)
                         0.00
                                0.07
                                       0.02
                                              0.10
                                                   0.00
                                                          0.04
                                0.37
                                       0.83
                                                                      Massive Py in Nodules
WB98, 477.50 #1
                         0.54
                                              0.61
                                                    0.27
                                                          0.22
                    (a)
                                0.05
                                       0.72
          duplicate (b)
                         0.44
                                              0.08
                                                    0.26
                                                          0.05
WB98, 477.50 #2
                    (a)
                         0.36
                                0.13
                                       0.53
                                              0.18 0.17
                                                          0.05
                                                                      Massive Py in Nodules
          duplicate (b)
                         0.46
                                0.05
                                       0.75
                                                   0.28
                                              80.0
                                                          0.05
                         0.10
                                0.13
                                       0.08
                                              0.18 -0.04 0.05
                                                                      Massive Py in Nodules
WB98, 477.50 #3
                    (a)
          duplicate (b)
                         0.05
                                0.05
                                       0.07
                                              0.08 0.00
                                                          0.05
WB98, 506.15
                         -0.68
                                0.08
                                       -0.93
                                                   -0.27
                                                          0.12
                                                                      Massive Py in Nodules
                    (a)
                                              0.11
                         -0.78
                                0.07
                                       -1.08
                                              0.10 -0.34 0.04
          duplicate (b)
                                0.23
                                       -1.27
                                              0.24 -0.40 0.02
                                                                      Euh. Py, disseminated in
WB98, 506.15 #1
                    (a)
                         -0.86
                                                                  0
                                                                      rock
          duplicate (b)
                         -0.73
                                0.05
                                       -1.06
                                              0.08 -0.35 0.05
WB98, 506.15 #2
                    (a)
                         -0.94
                                0.23
                                       -1.42
                                              0.24
                                                   -0.48 0.02
                                                                      Euh. Py, disseminated in
          duplicate (b)
                         -1.01
                                0.05
                                      -1.51
                                              0.08 -0.51 0.05
WB98, 506.15 #3
                         -0.87
                                0.06
                                       -1.26
                                              0.13 -0.39 0.09
                                                                      Euh. Pv. disseminated in
                    (a)
                                                                  0
                                                                      rock
          duplicate (b)
                         -0.78
                                0.05
                                      -1.13
                                              0.08 -0.37 0.05
WB98, 509.10
                         -2.41
                                       -3.52
                    (b)
                                0.07
                                              0.10 -1.15 0.04
                                                                      Massive Py in Nodules
                                                                  0
                         -2.75
                                0.06
                                       -4.09
WB98, 509.10 #1
                    (a)
                                              0.13 -1.33 0.09
                                                                      Nod py (2mm)
                                                                  0
                         -2.67
                                       -3.97
          duplicate (b)
                                0.05
                                              0.08 -1.31
                                                          0.05
WB98, 509.10 #2
                         -2.54
                                0.05
                                       -3.73
                                              0.11 -1.21
                                                                      Nod py (4mm)
                    (a)
                                                          0.07
          duplicate (b)
                         -2.48
                                0.05
                                       -3.61
                                              0.08 -1.15 0.05
WB98, 509.10 #3
                         -2.51
                                0.05
                                       -3.78
                                              0.11 -1.29 0.07
                                                                      Nod py (4mm)
                    (a)
          duplicate (b)
                         -2.44
                                0.05
                                       -3.64
                                              0.08 -1.22
                                                          0.05
                         -1.92
                                       -2.96
                                                                      Massive Py in Nodules
WB98, 513.60 a
                    (b)
                                0.07
                                              0.10 -1.03 0.04
                                                                  0
WB98, 513.60 b
                    (b)
                         -1.81
                                0.19
                                       -2.69
                                              0.25 -0.89 0.08
                                                                      Massive Py in Nodules
                                                                  0
          duplicate (c)
                         -1.83
                                0.07
                                       -2.77
                                              0.11
                         -2.13
                                0.28
                                       -3.19
                                                                      Massive Py in Nodules
WB98, 519.33
                    (a)
                                              0.34 -1.06 0.06
          duplicate (b)
                         -2.13
                                0.19
                                       -3.20
                                              0.25 -1.08 0.08
          duplicate (c)
                         -2.09
                                0.06
                                       -3.06
                                              0.05
WB98, 519.63 a
                         -1.28
                                0.28
                                       -1.92
                                              0.34 -0.65 0.06
                                                                      Massive Py in Nodules
                    (a)
          duplicate (b)
                         -1.22
                                0.19
                                       -1.84
                                              0.25 -0.63 0.08
          duplicate (c)
                         -1.22
                                0.12
                                       -1.92
                                              0.21
                                       -1.89
WB98, 519.63 b
                    (a)
                         -1.31
                                0.20
                                              0.32 -0.60 0.17
                                                                      Massive Py in Nodules
          duplicate (b)
                         -1.24
                                0.07
                                       -1.74
                                              0.10 -0.53
                                                          0.04
WB98, 519.63 #1
                    (a)
                         -1.36
                                0.24
                                       -2.04
                                              0.41 -0.68 0.16
                                                                      Nod py (1mm)
          duplicate (b)
                         -1.39
                                0.06
                                       -2.07
                                              0.09 -0.66 0.04
                                       -2.18
                         -1.44
                                0.05
                                              0.08 -0.75
                                                                      Nod py (1mm)
WB98, 519.63 #2
                    (a)
                                                          0.02
                                      -1.92
          duplicate (b)
                         -1.30
                                0.06
                                              0.09 -0.63 0.04
                                      -2.22
                                                                      Nod py (1mm)
WB98, 519.63 #3
                    (a)
                         -1.46
                                0.05
                                              0.08 -0.78 0.02
          duplicate (b)
                         -1.43
                                0.04
                                       -2.13
                                              0.06 -0.71
                                                          0.04
                                       -3.11
                                                                      Massive Py in Nodules
WB98, 520.85
                    (a)
                         -2.10
                                0.09
                                              0.17 -1.02 0.12
```

```
WB98, 520.85#1
                         -2.19
                                0.05
                                       -3.11
                                              0.07 -1.01 0.07
                                                                      Massive Py in Nodules
                    (a)
                         -2.20
                                       -3.20
           duplicate (b)
                                0.15
                                              0.24 -0.98
                                                          0.10
                         -1.98
                                0.05
                                       -2.85
                                              0.04 -0.89 0.02
                                                                      Massive Py in Nodules
WB98, 520.85#2
                    (a)
                                                                  0
          duplicate (b)
                         -2.03
                                0.15
                                       -2.93
                                              0.24 -0.92 0.10
                         -2.21
                                       -3.29
WB98, 520.85#3
                    (a)
                                0.13
                                              0.18 -1.06
                                                          0.07
                                                                  0
                                                                      Massive Py in Nodules
                         -2.15
                                0.08
                                       -3.32
                                                                      Massive Py in Nodules
WB98, 522.04
                    (b)
                                              0.12 -1.13 0.05
WB98, 523.55
                    (a)
                         -2.13
                                0.09
                                       -3.18
                                              0.08 -1.10 0.02
                                                                      Fine-g Py in nodules
           duplicate (b)
                         -2.24
                                0.15
                                       -3.32
                                              0.24 -1.11 0.10
ca. 2.6 Ga Carawine Dolomite, Hamersley Group, Hamersley Basin, Western Australia
                                       -5.18
                                              0.09 -1.68 0.04
RH2A-57, 148
                         -3.51
                                0.06
                                                                      Massive Pv
                    (a)
                         -3.39
                                0.08
                                       -4.97
                                              0.12 -1.60 0.05
          duplicate (b)
ca. 2.6 Ga Lokammona Formation, Schmidtsdrif Subgroup, Griqualand West Basin, South
Africa
SF-1, 599.8#1
                                0.13
                                       -4.92
                                              0.19 -1.63 0.07
                                                                      Massive Py in Nodules
                    (b)
                         -3.27
          duplicate (a)
                         -3.42
                                0.08
                                       -5.08
                                              0.13 -1.64 0.11
          duplicate (b)
                         -3.39
                                0.06
                                       -5.02
                                              0.08 -1.60 0.04
SF-1, 599.8#2
                    (b)
                         -3.32
                                0.13
                                       -4.87
                                              0.19 -1.58 0.07
                                                                      Massive Py in Nodules
                         -3.38
                                0.08
                                       -5.09
          duplicate (a)
                                              0.13 -1.70 0.11
                         -3.21
                                0.06
                                       -4.82
          duplicate (b)
                                              0.08 -1.59 0.04
                         -2.96
                                0.18
                                       -4.38
                                                                      Massive Py in Nodules
SF-1, 611.50
                    (a)
                                              0.39 -1.42 0.21
                         -2.98
                                0.19
                                       -4.37
                                              0.25 -1.40 0.08
          duplicate (b)
                                       -4.34
          duplicate (c)
                         -2.90
                                0.08
                                              0.12
SF-1, 611.50 #1
                         -3.04
                                0.08
                                       -4.50
                                              0.10 -1.48 0.02
                                                                      Nod py (1mm)
                    (a)
                         -2.93
                                       -4.36
          duplicate (b)
                                0.06
                                              0.09 -1.43 0.04
                         -2.90
                                0.08
                                       -4.31
SF-1, 611.50 #2
                    (a)
                                              0.10 -1.40 0.02
                                                                      Nod py (1mm)
          duplicate (b)
                         -2.87
                                0.04
                                       -4.30
                                              0.06 -1.40 0.04
                         -2.04
                                0.05
                                       -3.04
SF-1, 611.75
                    (a)
                                              0.08 -1.01 0.02
                                                                      Massive Py in Nodules
                                                                 +++
                                              0.10 -1.00 0.04
           duplicate (b)
                         -2.02
                                0.07
                                       -3.01
                                       -4.54
SF-1, 623.6
                    (a)
                         -3.05
                                0.05
                                              0.08 -1.50 0.03
                                                                      Fine-g & Massive Py in
                                                                  0
                                                                      nodules
          duplicate (b)
                         -2.96
                                0.07
                                       -4.40
                                              0.10 -1.44 0.04
                         -2.86
                                       -4.24
SF-1, 623.6 #1
                                0.05
                                              0.08 -1.34
                                                                      Massive Py in Nodules
                    (a)
                                                          0.02
                                0.06
                                       -4.13
          duplicate (b)
                         -2.77
                                              0.09 -1.33 0.04
SF-1, 623.6 #2
                    (a)
                         -2.80
                                0.05
                                       -4.13
                                              0.08 -1.33 0.02
                                                                      Massive Py in Nodules
                                                                  0
                         -2.79
                                       -4.15
          duplicate (b)
                                0.04
                                              0.06 -1.37
                                                          0.04
SF-1, 642.8
                         -0.07
                                0.14
                                       -0.06
                                              0.17 0.04
                                                                      Massive Py in Nodules
                    (a)
                                                          0.05
          duplicate (b)
                         -0.19
                                0.07
                                       -0.22
                                              0.10 0.00
                                                          0.04
                                                                      Euh py
SF-1, 642.8 #1
                         0.05
                                0.06
                                       0.01
                                              0.09 -0.02 0.04
                    (a)
                                       0.06
                         0.10
                                0.04
                                              0.06 -0.02 0.04
          duplicate (b)
SF-1, 642.8 #2
                    (a)
                         -0.04
                                0.05
                                       0.00
                                              0.08 0.04
                                                          0.04
                                                                      Euh py
                         0.00
          duplicate (b)
                                0.06
                                       0.03
                                              0.09
                                                    0.03
                                                          0.04
                                       0.10
SF-1, 2114.2
                    (a)
                          0.06
                                0.14
                                              0.17
                                                    0.08
                                                          0.05
                                                                      Fine-g Py
          duplicate (b)
                         0.15
                                0.07
                                       0.24
                                              0.10 0.13
                                                          0.04
          duplicate (b)
                         0.13
                                80.0
                                       0.24
                                              0.12 0.12
                                                          0.05
SF-1, 2175.5
                         0.22
                                0.05
                                       0.31
                                              0.08 0.14
                                                          0.02
                                                                      Fine-g Py
                    (a)
          duplicate (b)
                         0.21
                                0.07
                                       0.29
                                              0.10
                                                    0.13
                                                          0.04
                                       0.22
          duplicate (a)
                         0.13
                                0.09
                                              0.15 0.08 0.07
                                0.08
                                       0.33
          duplicate (b)
                         0.22
                                              0.12 0.10 0.05
ca. 2.65 Ga Lewin Shale / Jeerinah Formation, Upper part of the Fortescue Group, Hamersley
Basin, Western Australia
RHDH 2A, 280.26
                                0.14
                                       -1.12 0.14 -0.32 0.05
                                                                      Massive Py
                         -0.82
                         -0.69
                                      -0.92 0.08 -0.24 0.04
          duplicate (b)
                                0.06
```

2.63 Ga Roy Hill Shale Member of the Jeerinah Formation, Upper part of the Fortescue Group, Hamersley Basin, Western Australia									
FVG-1, 707.95	(b)	-1.27	0.33	-1.91	0.43	-0.53	0.08	+	Fine-g Py in nodules
duplicate	1. 1	-1.26	0.19	-1.87	0.20		0.10		,
FVG-1, 707.95 #1	(a)	-1.68	0.05	-2.49	0.09			+	Nod py (~cm)
duplicate	٠,	-1.69	0.05	-2.49					7 ()
FVG-1, 707.95 #2	(a)	-1.41	0.05	-2.14		-0.72		+	Nod py (~cm)
duplicate	(b)	-1.42	0.05	-2.15		-0.72			17()
FVG-1, 722.6 #1	(a)	-1.94	0.06	-2.84	0.09	-0.92	0.02	++	Massive Py in Nodules
duplicate	(b)	-1.93	0.05	-2.77	0.08	-0.87	0.05		•
FVG-1, 722.6 #2	(a)	-1.94	0.08	-2.97	0.10	-1.03	0.04	++	Massive Py in Nodules
duplicate	(b)	-2.02	0.05	-3.04	0.08	-1.03	0.05		·
FVG-1, 751.77	(a)	-2.36	0.10	-3.58	0.12	-1.22	0.03	++	Massive Py in Nodules
duplicate	(b)	-2.45	0.19	-3.73	0.25	-1.28	0.08		·
duplicate	(c)	-2.34	0.09	-3.56	0.09				Hand-picked pyrite
									nodules
FVG-1, 752.8	(a)	-2.78	0.06	-4.11		-1.32		+	Massive Py in Nodules
duplicate	(b)	-2.72	0.07	-4.01		-1.30			
FVG-1, 752.8 #1	(a)	-3.06	0.07	-4.54		-1.47		+	Nod py (~cm)
duplicate		-3.08	0.06	-4.58		-1.48			
FVG-1, 752.8 #2	(a)	-2.58	0.07	-3.86		-1.26	0.10	+	Nod py (~cm)
duplicate	(b)	-2.55	0.06	-3.84			0.04		
FVG-1, 761.8	(a)	-1.09	0.05	-1.89		-0.73		0	Massive Py in Nodules
duplicate	٠,	-1.13	0.19	-1.85		-0.71	0.10		
FVG-1, 761.8 #1	(a)	-1.17	0.05	-1.70	0.15	-0.52		0	Euh py
duplicate		-1.11	0.06	-1.65	0.09	-0.53			
FVG-1, 761.8 #2	(a)	-1.22	0.05	-1.74	0.15	-0.53	0.10	0	Euh py
duplicate	(b)	-1.14	0.06	-1.66	0.09	-0.53	0.04		
FVG-1, 777.8	(a)	-1.37	0.19	-1.91	0.22			0	Euh py in nodules
duplicate	` '	-1.38	0.07	-1.95	0.10	-0.56	0.04		
FVG-1, 781.80 a	(a)	-1.91	0.19	-2.74		-0.88	0.00	0	Euh py in nodules
duplicate	٠,	-1.92	0.07	-2.78	0.10		0.04		E to a facility
FVG-1, 781.80 b	(a)	-2.08	0.18	-3.08	0.39	-1.01	0.21	0	Euh py in nodules
duplicate	٠,	-2.14	0.19	-3.13	0.25	-1.00	0.08		
duplicate	(c)	-2.10	0.07	-3.12	0.11	0.77	0.40		Fub ny in nadulaa
FVG-1, 784.1	(a)	-1.74		-2.52				+	Euh py in nodules
duplicate FVG-1, 787.4	. ,	-1.80 -1.50	0.16 0.06	-2.63 -2.11		-0.83 -0.75			Massive Py in nodules
·	(a)	-1.30 -1.45	0.06	-2.11 -2.22		-0.73		+	Massive Fy III Hoddies
duplicate FVG-1, 791.55	(a) (a)	-1.45	0.09	-1.61	0.12			+	Euh py in nodules
duplicate	٠,,	-1.13	0.03	-1.51		-0.42		т	Luii py iii fiodules
FVG-1, 799.0 #1	(a)	-0.20	0.06	-0.34		-0.42		+	Massive Py in nodules
duplicate	. ,	-0.20 -0.17	0.06	-0.3 4 -0.28		-0.13		т	Massive Fy III Houdies
FVG-1, 799.0 #2	(a)	-0.76	0.05	-1.21		-0.10		+	Massive Py in nodules
duplicate	` '	-0.74	0.06	-1.18		-0.42		т	Massive i y ili lloddies
FVG-1, 814.75	(a)	-1.73	0.05	-2.66		-0.42		0	Euh py in nodules
duplicate	. ,	-1.62	0.03	-2.47		-0.81		J	Early in Hoddies
FVG-1, 815.9	(a)	-1.56	0.07	-2.35		-0.81		0	Massive & Fine-g Py in
1 10 1,010.0	(α)	1.50	0.00	2.00	0.10	0.01	0.12	J	nodules
duplicate	(b)	-1.42	0.33	-2.10	0.43	-0.66	0.08		
FVG-1, 826.55	(a)	-1.48	0.05	-2.17		-0.67		0	Massive & Euh Py in
	•								nodules

```
duplicate (b)
                       -1.48
                               0.07
                                     -2.13
                                            0.10 -0.64 0.04
FVG-1, 830.6
                        -1.28
                               0.25
                                     -1.76
                                             0.33 -0.51 0.13
                    (a)
                                                                    Fine-q Py in nodules
          duplicate (b)
                        -1.33
                               0.19
                                     -1.94
                                             0.20 -0.60 0.10
FVG-1, 830.6 #1
                    (a)
                        -1.70
                               0.06
                                     -2.51
                                             0.09 -0.82 0.02
                                                                    Nod py (~cm)
                                                               ++
                                     -2.38
                        -1.62
                               0.06
                                             0.09 -0.77 0.04
          duplicate (b)
FVG-1. 830.6 #2
                        -1.87
                               0.05
                                     -2.82
                                            0.08 -0.95 0.02
                                                                    Nod py (~cm)
                    (a)
          duplicate (b)
                        -1.90
                               0.04
                                     -2.86
                                            0.06 -0.96 0.04
FVG-1, 835.55
                        -1.66
                               0.07
                                     -2.36
                                             0.11 -0.69 0.02
                                                                    Massive & Euh Py in
                    (a)
                                                                0
                                                                    nodules
          duplicate (b)
                        -1.74
                               0.07
                                     -2.48
                                             0.10 -0.74 0.04
FVG-1, 849.6
                        -0.55
                               0.07
                                     -0.89
                                            0.10 -0.33 0.04
                    (b)
                                                                    Massive & Fine-q Py in
                                                                0
                                                                    nodules
          duplicate (b)
                        -0.62
                               0.19
                                     -1.00
                                            0.19 -0.36 0.10
FVG-1, 849.6 #1
                        -0.95
                               0.05
                                     -1.40
                                            0.13 -0.44 0.10
                    (a)
                                                                    Euh py in nodules
          duplicate (b)
                        -0.97
                               0.05
                                     -1.46
                                            0.06 -0.47 0.04
FVG-1, 849.6 #2
                    (a)
                        -0.87
                               0.08
                                     -1.26
                                            0.13 -0.40 0.10
                                                                    Euh py in nodules
          duplicate (b) -0.90 0.05
                                    -1.33
                                            0.06 -0.44 0.04
ca. 2.72 Ga Helen Iron Formation, Michipicoten Group, Sir James Mines, Wawa, ON, Canada
WW1
                    (b) -1.76
                               0.16
                                    -2.64 0.13 -0.88 0.09
                                                                    nodular Py
ca. 2.72 Ga Deer Lake greenstone sequence, MN, USA
#26503. 1000'
                        -1.42
                               0.11
                                     -2.02
                                            0.12 -0.64 0.02
                                                                    Massive Py
                    (a)
          duplicate (b)
                        -1.31
                               0.15
                                     -1.82
                                            0.24 -0.55 0.10
                                                                    Massive & Euh Py
#26503, 1095'
                    (a)
                        -1.93
                               0.07
                                     -2.91
                                            0.11 -0.98 0.00
                                                                0
          duplicate (b)
                        -1.85
                               0.07
                                     -2.78
                                            0.10 -0.95 0.04
                        -1.90
                                     -2.79
                                            0.20 -0.90 0.11
#26503, 1117'
                    (a)
                               0.10
                                                                    Massive Py
#26516, 366'
                    (b)
                        -1.94
                               0.15
                                     -2.93
                                            0.24 -1.01 0.10
                                                                    Massive Py
                                                                0
                        -1.87
                               0.06
                                     -2.87
                                             0.08 -0.99 0.04
          duplicate (b)
                               0.08
                                     -2.26
                                            0.11 -0.84 0.13
#26516, 378'
                    (a)
                        -1.43
                                                                    Massive & Euh Py
          duplicate (b) -1.37
                               0.07
                                     -2.14
                                            0.10 -0.78 0.04
```

(a): Instrumental mass bias corrected using the "standard-sample bracketing" method; (b): Instrumental mass bias corrected using Nickel standard solution as an internal standard (c) analysis performed on the Nu plasma MC-ICPMS at the University of Cambridge. Sample descripton: Py: pyrite; Fine-g: fine grained crystals; Euh.: euhedral minerals: Nod: nodules (round shape and concentric banding). Org. C: visual estimate of organic carbon content associated with pyrite grains: (o) no org. C (+) trace org. C (++) minor org. C (<1%) (+++) major org. C (<10%).

Table S3: Fe-isotope	composition	of Iron	Formations
I able 33. Fe-18010bi		oi iioii	FUIIIIaliUlis

Sample	Meth	δ^{56} Fe	2σ	δ^{57} Fe	2σ	$\delta^{57/56}$	2σ	Description			
	od					Fe					
Upper Cherty Member, ca. 1.88 Ga Biwabik Iron Formation, Cliffs Erie Mine, Mesabi Range, MN USA											
Biwabik #hem	(a)	-0.07	0.05	-0.04	0.08	0.04	0.02	Massive Hem within silica matrix			
duplicate	(b)	-0.07	0.16	0.01	0.13	0.02	0.09				
Biwabik #mag	(b)	-0.23	0.08	-0.31	0.12	-0.09	0.05	Massive Mag within silica matrix			
duplicate	(b)	-0.24	0.16	-0.38	0.13	-0.11	0.09				
ca. 1.88 Ga Ironwood Iron Formation, Tyler Forks River, WI, USA											
4TyF-10 #mag	(a)	0.48	0.17	0.58	0.29	0.13	0.10	Massive Mag within silica matrix			
duplicate		0.45	0.06	0.57	0.08	0.13	0.04				
ca. 2.47 Ga Kuru	ıman I	ron For	matio	n, Kurur	nan are	ea, Gric	qualan	d West Basin, South Africa			
KU-9-537 #mag	(a)	0.46	0.05	0.77	0.08	0.26	0.02	Massive Mag within silica matrix			
duplicate	(b)	0.45	0.08	0.77	0.12	0.27	0.05				
duplicate	(b)	0.37	0.16	0.57	0.13	0.17	0.09				
ca. 2.47 Ga Brockman Iron Formation, Wittenoom Gorge, Western Australia											
WIT-15 #mag	(a)	0.65	0.08	0.99	0.08	0.33	0.04	Very Fine-g Mag			
duplicate	(b)	0.62	0.08	0.93	0.12	0.30	0.05				
WIT-46 #mag	(b)	0.25	0.08	0.41	0.12	0.15	0.05	Very Fine-g Mag			
ca. 2.7 Ga Manje	ca. 2.7 Ga Manjeri Formation, Belingwe Belt, Zimbabwe (Rouxel, Bickle & Galy, unpub data)										
MHZ 7b	(c)	1.02	0.08	1.48	0.03	n.d.		Fe-oxides (magnetite; hematite)			
MHZ 7e	(c)	1.06	0.13	1.52	0.05	n.d.		Fe-oxides (magnetite; hematite)			
TR 40	(c)	1.18	0.06	1.68	0.09	n.d.		Fe-oxides (magnetite; hematite)			
TR 42	(c)	1.16	0.13	1.70	0.18	n.d.		Fe-oxides (magnetite; hematite)			
TR 43	(c)	1.61	0.13	2.47	0.13	n.d.		Fe-oxides (magnetite; hematite)			
ca. 2.72 Ga Hele	ca. 2.72 Ga Helen Iron Formation, Michipicoten Group, MacLeod Mine, Wawa, Ontario, Canada										
Stromato-carb	(b)	-1.28	0.06	-1.90	0.08	-0.60	0.04	honey-colored carbonate			
duplicate	(b)	-1.33	0.16	-1.96	0.13	-0.65	0.09	•			
Stromato-mag	(b)	-0.75	0.06	-1.05	0.08	-0.31	0.04	Fine-g Euh mag			
duplicate	(a)	-0.71	0.20	-1.09	0.43	-0.37	0.22				
duplicate		-0.71	0.16	-1.07	0.13	-0.38	0.09				
Stromato-py	(a)	-0.91	0.09	-1.35	0.20	-0.41	0.13	Euhedral Py crystals			
duplicate	(b)	-1.01	0.06	-1.53	0.08	-0.52	0.04				

(a): Instrumental mass bias corrected using the "standard-sample bracketing" method; (b): Instrumental mass bias corrected using Nickel standard solution as an internal standard (c) analysis performed on the Nu plasma MC-ICPMS at the University of Cambridge. Hem: hematite; Mag: magnetite; stromato: stromatolite facies composed mainly of Fe-rich carbonate (carb), disseminated fine grained magnetite and euhedral pyrite. Fine-g: fine grained crystals; Euh.: euhedral minerals.

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