# Multi-frame Full Matrix Capture (MFMC) File Format Specification

# Version 1.0.0

## Summary

The Multi-frame Full Matrix Capture (MFMC) file format is designed for storage of data obtained from ultrasonic array probes. It has been designed specifically for the storage of the original ultrasonic data in its raw form acquired under the paradigm of Full Matrix Capture (FMC), although the format can accommodate other acquisition modalities.

The quantum of data in an MFMC file is a single frame of FMC data. A collection of frames where all acquisition parameters except the probe position are fixed constitutes a sequence. Typically, a sequence represents the results of a probe being scanned over a target. One MFMC file can accommodate multiple sequences. The number of frames in any sequence is limited only by physical storage capacity and operating system constraints. Frames can be appended to sequences within an existing file.

The underlying file format is the Hierarchical Data Format (HDF5), which is structured as a hierarchy of groups containing data-fields. A HDF5 file is a valid MFMC file, if the groups and data-fields defined by the MFMC specification are present and conform to the MFMC specification. A valid MFMC file can contain an unlimited number of additional groups and data-fields to provide additional functionality to satisfy the requirements of users.

## Introduction

\*Give the context and include at least one schematic of a typical use case (array being scanned over component with multiple frames in sequence).

\*use cases

Fig.  Example use cases: (a) single probe scanned over component; (b) pair of probes scanned over component.

## Definitions

* Element – an individually-addressable ultrasonic transduction device;
* Placement – position and orientation of an entity (e.g. an element);
* Probe – a collection of elements with placements that are fixed relative to one another;
* Probe Element Combination (PEC) – the double index system used within an MFMC file to identify a specific element in a specific probe;
* Focal law – a set of instructions that specify how one or more PECs are used together;
* Transmit focal law – a focal law relating to transmission of ultrasound from one or more PECs;
* Receive focal law – a focal law relating to reception of ultrasound from one or more PECs;
* Acquisition parameters – the collection of parameters (including transmit and receive focal laws) that specify how ultrasonic data is acquired;
* A-scan – a time-domain, un-rectified ultrasonic signal (comprising amplitude measurements regularly sampled in time at a specified sampling frequency) that is recorded for a combination of transmit focal law and receive focal law;
* Frame – a collection of A-scans obtained using different transmit and receive focal laws for each A-scan;
* Sequence – a collection of frames in which all acquisition parameters except the probe position are fixed from one frame to another;
* Full Matrix Capture (FMC) – a frame of data from PECs in which the transmit and receive focal laws each specify a single PEC and A-scans from all possible combinations of transmit and receive focal laws are recorded;
* Half Matrix Capture (HMC) – a subset of FMC where acoustic reciprocity is assumed and A-scans from only the unique transmit and receive focal law combinations are recorded;
* Plane Wave Imaging (PWI) – an alternative data-acquisition modality where multiple elements are used in transmission to emit a plane wave at a specified angle;
* Probe Coordinate System (PCS) – coordinate systems (one for each probe) that describe the (fixed) relative placement of elements within a probe;
* Global Coordinate System (GCS) – the single fixed laboratory coordinate system;
* Ultrasonic time – timescale over which an individual ultrasonic A-scan is recorded, which is assumed to be instantaneous compared to timescale associated with mechanical movement of probes;

## MFMC File Structure

\*explanation of template. Common points: e.g. units, character sets, byte ordering, row-major etc

### Overview and Naming Conventions

The overall structure of an MFMC file is summarised in Fig. 1.

Fig. 1 Overall file structure. User-specified groups and data-fields can be added as necessary at any level.

A HDF5 file contains a hierarchy of groups that contain datasets and attributes. The difference between datasets and attributes is that the latter are intended for small, fixed-size data, while the former are designed for holding multi-dimensional arrays of data. In all other respects their behaviour is the same and for brevity, the term datafield will be used to describe both unless stated.

Groups, datasets and attributes within the hierarchy are accessed via paths with the different levels in the hierarchy indicated by the / character. In the MFMC file, paths are capitalised, e.g.

* /GROUP\_LEVEL\_1/…/GROUP\_LEVEL\_N/DATASET\_NAME – dataset;
* /GROUP\_LEVEL\_1/…/GROUP\_LEVEL\_N/ATTRIBUTE\_NAME – attribute.

On occasions, multiple instances of the same type of group (e.g. probes) are needed at the same level in the hierarchy. These are denoted by positive, integer, numerical values appended to the group name in angle brackets, e.g. /PROBE<1>. Note that the numerical values are not numerical indices in the programmatical sense; they are just part of the sequence of characters in the path.

The notation used in this document for datafield paths is as follows

* /A\_{B/C} – shorthand for specific paths /A\_B and /A\_C;
* /A\_\* – shorthand for any paths that match pattern with \* as wildcard, e.g. /A\_B and /A\_C;
* B in /A – shorthand for path /A/B;
* /A<x> – placeholder for path indexed by numerical value x, e.g. /A<1>, /A<2> etc.

### Dimensions, Sizes and Cross-referencing

Datafields in the MFMC file are typically multi-dimensional arrays of numbers. The size of a datafield is given in square brackets with dimension separated by commas, e.g. [2,3]. Therefore [1] represents a single scalar quantity, [5] represents a vector and [2,3] represents a matrix.

Three types of dimension size can be identified:

* specific fixed value – e.g. the size of the first dimension of /PROBE<p>/ELEMENT\_POSITION is always 3 as it represents coordinates in the PCS;
* arbitrary fixed value – these are fixed at creation to a specific value and may also have to be consistent with the size of dimensions in other datafields, e.g. the size of the second dimension of the three datafields /PROBE<p>/ELEMENT\_{POSITION/MINOR/MAJOR} must equal the number of elements in the probe;
* expandable – the size of these dimensions is not fixed at creation and typically represent quantities that can be appended with more data as it is acquired, e.g. /MFMC<m>/MFMC\_DATA.

Dimension sizes that are variable or expandable are identified by the variable names listed in Table 1.

|  |  |
| --- | --- |
| **Variable** | **Description** |
| N\_P | Number of probes (expandable) |
| N\_M | Number of MFMC sequences in file (expandable) |
| N\_E<p> | Number of elements in p-th probe |
| N\_F<m> | Number of FMC frames in m-th sequence of MFMC data (expandable) |
| N\_T<m> | Number of time points per A-scan in m-th sequence |
| N\_A<m> | Number of A-scans per frame in m-th sequence |
| N\_B<m> | Number of firing events in m-th sequence |
| N\_Q<m> | Number of probes used in m-th sequence |
| N\_L<m> | Number of focal laws associated with each frame in m-th sequence |
| N\_C<m><k>] | Number of probe/element combinations used in k-th focal law in m-th sequence |
| P | Probe number |
| M | MFMC sequence number |
| K | Focal law number |

Table 1 List of variables used in file structure description.

Some datafields contain integers that are cross-references to the indices of other entities. For example, in /MFMC<m>/COMMON the integers in TRANSMIT\_LAW\_INDEX refer to the indices of focal laws and the integers in LAW<k>/ELEMENT refer to the indices of the elements in a probe used in a focal law. Here two types of indices can be identified:

* indices to paths of groups – e.g. TRANSMIT\_LAW\_INDEX in /MFMC<m>/COMMON refers to the numbers k in LAW<k>;
* indices into dimensions of datafields – e.g. LAW<k>/ELEMENT in /MFMC<m>/COMMON indexes into the second dimension of ELEMENT\_{POSITION/MINOR/MAJOR} in /PROBE<p>.

For an MFMC file to be valid, cross-referencing datafields must refer to valid indices, which means:

* paths of groups cross-referenced by indices must exist in the file;
* indices into a datafield dimension of size N must be in the range 1...N.

### Detailed Structure

Table 2 provides the complete list of the datafields in the MFMC file specification.

| **Path** | **M or O** | **D or A** | **Class** | **Dims** | **Size** |
| --- | --- | --- | --- | --- | --- |
| /VERSION | M | A | H5T\_STRING | 1 | [1] |
| /PROBE<p>/ELEMENT\_POSITION | M | D | H5T\_FLOAT | 2 | [3,N\_E<p>] |
| /PROBE<p>/ELEMENT\_MINOR | M | D | H5T\_FLOAT | 2 | [3,N\_E<p>] |
| /PROBE<p>/ELEMENT\_MAJOR | M | D | H5T\_FLOAT | 2 | [3,N\_E<p>] |
| /PROBE<p>/ELEMENT\_SHAPE | M | D | H5T\_INTEGER | 1 | [N\_E<p>] |
| /PROBE<p>/ELEMENT\_RADIUS\_OF\_CURVATURE | O | D | H5T\_FLOAT | 1 | [N\_E<p>] |
| /PROBE<p>/ELEMENT\_AXIS\_OF\_CURVATURE | O | D | H5T\_FLOAT | 2 | [3,N\_E<p>] |
| /PROBE<p>/WEDGE\_SURFACE\_POINT | O | A | H5T\_FLOAT | 1 | [3] |
| /PROBE<p>/WEDGE\_SURFACE\_NORMAL | O | A | H5T\_FLOAT | 1 | [3] |
| /PROBE<p>/DEAD\_ELEMENT | O | D | H5T\_INTEGER | 1 | [N\_E<p>] |
| /PROBE<p>/CENTRE\_FREQUENCY | M | A | H5T\_FLOAT | 1 | [1] |
| /PROBE<p>/BANDWIDTH | O | A | H5T\_FLOAT | 1 | [1] |
| /PROBE<p>/PROBE\_MANUFACTURER | O | A | H5T\_STRING | 1 | [1] |
| /PROBE<p>/PROBE\_SERIAL\_NUMBER | O | A | H5T\_STRING | 1 | [1] |
| /PROBE<p>/PROBE\_TAG | O | A | H5T\_STRING | 1 | [1] |
| /PROBE<p>/WEDGE\_MANUFACTURER | O | A | H5T\_STRING | 1 | [1] |
| /PROBE<p>/WEDGE\_SERIAL\_NUMBER | O | A | H5T\_STRING | 1 | [1] |
| /PROBE<p>/WEDGE\_TAG | O | A | H5T\_STRING | 1 | [1] |
| /MFMC<m>/MFMC\_DATA | M | D | H5T\_FLOAT / H5T\_INTEGER | 3 | [N\_T<m>,N\_A<m>,N\_F<m>] |
| /MFMC<m>/MFMC\_DATA\_IM | O | D | H5T\_FLOAT / H5T\_INTEGER | 3 | [N\_T<m>,N\_A<m>,N\_F<m>] |
| /MFMC<m>/PROBE\_PLACEMENT\_INDEX | M | D | H5T\_INTEGER | 2 | [N\_A<m>,N\_F<m>] |
| /MFMC<m>/PROBE\_POSITION | M | D | H5T\_FLOAT | 3 | [3,N\_Q<m>,N\_B<m>] |
| /MFMC<m>/PROBE\_X\_DIRECTION | M | D | H5T\_FLOAT | 3 | [3,N\_Q<m>,N\_B<m>] |
| /MFMC<m>/PROBE\_Y\_DIRECTION | M | D | H5T\_FLOAT | 3 | [3,N\_Q<m>,N\_B<m>] |
| /MFMC<m>/COMMON/TRANSMIT\_LAW\_INDEX | M | D | H5T\_INTEGER | 1 | [N\_A<m>] |
| /MFMC<m>/COMMON/RECEIVE\_LAW\_INDEX | M | D | H5T\_INTEGER | 1 | [N\_A<m>] |
| /MFMC<m>/COMMON/PROBE\_INDEX | M | D | H5T\_INTEGER | 1 | [N\_P<m>] |
| /MFMC<m>/COMMON/TIME\_STEP | M | A | H5T\_FLOAT | 1 | [1] |
| /MFMC<m>/COMMON/START\_TIME | M | A | H5T\_FLOAT | 1 | [1] |
| /MFMC<m>/COMMON/SPECIMEN\_VELOCITY | M | A | H5T\_FLOAT | 1 | [2] |
| /MFMC<m>/COMMON/WEDGE\_VELOCITY | O | A | H5T\_FLOAT | 1 | [2] |
| /MFMC<m>/COMMON/TAG | O | A | H5T\_STRING | 1 | [1] |
| /MFMC<m>/COMMON/DAC\_CURVE | O | D | H5T\_FLOAT | 1 | [N\_T<m>] |
| /MFMC<m>/COMMON/RECEIVER\_AMPLIFIER\_GAIN | O | A | H5T\_FLOAT | 1 | [1] |
| /MFMC<m>/COMMON/FILTER\_TYPE | O | A | H5T\_INTEGER | 1 | [1] |
| /MFMC<m>/COMMON/FILTER\_PARAMETERS | O | A | H5T\_FLOAT | 2 | [2,N\_F<m>] |
| /MFMC<m>/COMMON/FILTER\_DESCRIPTION | O | A | H5T\_STRING | 1 | [1] |
| /MFMC<m>/COMMON/OPERATOR | O | A | H5T\_STRING | 1 | [1] |
| /MFMC<m>/COMMON/DATE\_AND\_TIME | O | A | H5T\_STRING | 1 | [1] |
| /MFMC<m>/COMMON/LAW<k>/PROBE | M | D | H5T\_INTEGER | 1 | [N\_C<m><k>] |
| /MFMC<m>/COMMON/LAW<k>/ELEMENT | M | D | H5T\_INTEGER | 1 | [N\_C<m><k>] |
| /MFMC<m>/COMMON/LAW<k>/DELAY | O | D | H5T\_FLOAT | 1 | [N\_C<m><k>] |
| /MFMC<m>/COMMON/LAW<k>/WEIGHTING | O | D | H5T\_FLOAT | 1 | [N\_C<m><k>] |

Table 2 List of the datafields in an MFMC file. Abbreviations: M/O = Mandatory / Optional; D/A = Dataset / Attribute; PCS = Probe Coordinate System; GCS = Global Coordinate System. Mandatory / Optional is applicable to children if parent dataset exists, i.e. a mandatory parameter in an optional group is mandatory if the group exists.

### Validity of MFMC File

An MFMC is valid if the following requirements are satisfied:

* all mandatory datafields listed in Table 2 are present;
* the class of data in datafields listed in Table 2 (whether mandatory or optional) is as specified;
* the number of dimensions of datafields listed in Table 2 (whether mandatory or optional) is as specified;
* fixed dimension sizes of datafields listed in Table 2 (whether mandatory or optional) is as specified;
* variable dimension sizes of datafields listed in Table 2 (whether mandatory or optional) are self-consistent;
* cross-referencing datafields refer to valid indices.

## Explanatory Notes

### Version (/VERSION)

The MFMC version is given by a string in the form MAJOR.MINOR.PATCH (e.g. 1.0.0) according to the Semantic Versioning 2.0.0 convention [<https://semver.org/>], where {MAJOR/MINOR/PATCH} are all integer numbers without leading zeros. For development purposes the string may be appended with further information following a hyphen, e.g. 2.3.17-beta. The interpretation of the three numerical values is:

* MAJOR – for changes that require a new Application Programming Interface (API);
* MINOR – for added functionality that is backwards-compatible with existing API;
* PATCH – for bug-fixes that are backwards-compatible with existing API.

### Probe (/PROBE<p>)

A probe group specifies an ultrasonic array that contains multiple elements. An MFMC file can contain multiple probe definitions, which are enumerated by the index <p> where p is an integer starting at 1 without leading zeros. Note that <p> is part of the string defining the name of the group in the underlying HDF5 file; it is not an index in the programmatical sense.

The defining characteristic of a probe is that the spatial location of the elements within one probe are fixed relative to each other. The positions and orientations of elements are defined in terms of a local Probe Coordinate System (PCS) for each probe. The PCS is a right-handed Cartesian system described by the axes . The recommended convention for axis orientation for array probes with elements in a single plane is as follows:

* -axis – perpendicular to plane of elements, origin in the plane of elements, positive direction orientated in direction of ultrasonic emission, as shown in Fig. 2(a);
* -axis – in plane of elements, aligned to major dimension of probe, orientated in direction of increasing element index for a 1-D linear probe or in direction of most-rapidly increasing element index in the case of a 2D matrix array probe, as shown in Fig. 2(b);
* -axis – in plane of elements, aligned to minor dimension of probe, orientated in direction of least-rapidly increasing element index in the case of a 2D matrix array probe, as shown in Fig. 2(c).

\*PCS conventions\*

Fig. 2 Recommended conventions for Probe Coordinate System (PCS) for: (a) general case; (b) 1D linear probe; (c) 2D matrix array probe.

#### Array Elements (/PROBE<p>/ELEMENT\_\*)

Within /PROBE<p>, the placement of each element in the PCS is given by the three [3,N\_E<p>] datafields ELEMENT\_POSITION and ELEMENT\_{MAJOR/MINOR}. For all three element-placement datafields, the first dimension represents the components of a vector in the PCS and the second dimension represents the element index within the probe. The centre of each element is defined by the position vector, , in ELEMENT\_POSITION as shown in Fig. 3. The orientation of each element is defined by the vectors in ELEMENT\_MAJOR and ELEMENT\_MINOR. For each element these are vectors, and , from the centre of the respective element to one of the tips of its major and minor axes respectively. The vectors and for each element should be orthogonal and in the plane of the element. If they are not orthogonal then should be given priority and the component of that is orthogonal to should be used for purposes such as determining the width of a rectangular element. The vectors and should be orientated such that their cross-product, , is in the direction of ultrasonic emission. The shape of an element is denoted by the integer value in the array ELEMENT\_SHAPE. Currently only rectangular and elliptical elements are supported according to:

* 1 – rectangular, dimensions by , as shown in Fig. 3(a);
* 2 – elliptical, major axis length , minor axis length , as shown in Fig. 3(b).

Focused elements are specified through the optional ELEMENT\_RADIUS\_OF\_CURVATURE and ELEMENT\_AXIS\_OF\_CURVATURE datafields. These are interpreted as follows:

* Neither ELEMENT\_RADIUS\_OF\_CURVATURE or ELEMENT\_AXIS\_OF\_CURVATURE exists – elements are planar;
* ELEMENT\_RADIUS\_OF\_CURVATURE exists but ELEMENT\_AXIS\_OF\_CURVATURE does not – each element is a portion of a sphere with centre at a distance of ELEMENT\_RADIUS\_OF\_CURVATURE in the direction relative to ELEMENT\_POSITION, as shown in Fig. 3(c);
* ELEMENT\_RADIUS\_OF\_CURVATURE and ELEMENT\_AXIS\_OF\_CURVATURE both exist – each element is a portion of a cylinder with axis in the direction ELEMENT\_AXIS\_OF\_CURVATURE passing through a point at a distance of ELEMENT\_RADIUS\_OF\_CURVATURE in the direction relative to ELEMENT\_POSITION, as shown in Fig. 3(d).

\*Element position and orientation vector schematic

Fig. 3 Element geometry description in PCS: (a) rectangular element; (b) elliptical element; (c) spherically-focused element; (d) cylindrically-focused element.

#### Wedge (/PROBE<p>/WEDGE\_SURFACE\_{POINT/NORMAL})

These optional parameters can be used to specify the geometry of a (rigid) coupling wedge with a planar working surface. The [3] attribute WEDGE\_SURFACE\_POINT specifies the coordinates (in the PCS) of a point on the working surface of the wedge and the [3] attribute WEDGE\_SURFACE\_NORMAL specifies the working surface normal vector, as shown in Fig. 3.

\*Wedge

Fig. 3 Wedge geometry description in PCS.

#### Other Probe Datafields

\*Todo for

/PROBE<p>/DEAD\_ELEMENT

/PROBE<p>/CENTRE\_FREQUENCY

/PROBE<p>/BANDWIDTH

/PROBE<p>/PROBE\_MANUFACTURER

/PROBE<p>/PROBE\_SERIAL\_NUMBER

/PROBE<p>/PROBE\_TAG

/PROBE<p>/WEDGE\_MANUFACTURER

/PROBE<p>/WEDGE\_SERIAL\_NUMBER

/PROBE<p>/WEDGE\_TAG

### MFMC Sequence (/MFMC<m>)

An MFMC sequence contains one or more frames of data acquired under identical conditions except for probe position(s). The group /MFMC<m>/COMMON contains all the information which is common to the sequence. The actual data is in the expandable [N\_T<m>,N\_A<m>,N\_F<m>] dataset /MFMC<m>/MFMC\_DATA (and for complex data the optional [N\_T<m>,N\_A<m>,N\_F<m>] dataset /MFMC<m>/MFMC\_DATA\_IM may be used to record the imaginary component). The datafields /MFMC<m>/PROBE\_POSITION and /MFMC<m>/PROBE\_{X/Y}\_DIRECTION describe probe placement; each A-scan in each frame is cross-referenced to one probe placement in these datafields.

#### Focal Laws (/MFMC<m>/COMMON/LAW<k>)

Each A-scan in a frame must be associated with both a transmit and a receive focal law through the datafields /MFMC<m>/COMMON/{TRANSMIT/RECEIVE}\_LAW\_INDEX (note that these apply to all the frames in a sequence). These provide the index k into the groups /MFMC<m>/COMMON/LAW<k> which provide the detailed description of each focal laws(the same structure is used for both transmit and receive focal laws). In /MFMC<m>/COMMON/LAW<k>, the mandatory datafields are PROBE and ELEMENT. These indicate the PECs used for the focal law. For FMC/HMC data, only one PEC is used in transmission and one PEC in reception, so these datafields contain single values; however, for other acquisition modalities such as PWI, a focal law may specify multiple PECs. The optional datafields, DELAY and WEIGHTING in /MFMC<m>/COMMON/LAW<k>, specify the relative delay (in ultrasonic time) and weighting between the different PECs in the focal law. The default values if these datafields are not present are zero delay and unity weighting for all PECs.

#### Timebase (/MFMC<m>/COMMON/TIME\_STEP and /MFMC<m>/COMMON/START\_TIME)

All A-scans in all frames in a sequence must be uniformly sampled on a common ultrasonic time axis. Ultrasonic time zero on this axis should correspond to the instant when an ultrasonic pulse is emitted from the probe (for FMC/HMC data this is unambiguous, but for focal laws involving delays, the user needs to define exactly what time zero corresponds to). The time axis parameters are specified in /MFMC<m>/COMMON/by the scalar attributes:

* START\_TIME – the ultrasonic time of the first point in the A-scan (which need not be zero if, for example, the early portion of a signal is not of interest);
* TIME\_STEP – the time step between adjacent points.

#### Ultrasonic Velocities (/MFMC<m>/COMMON/{SPECIMEN/WEDGE}\_VELOCITY)

Ultrasonic velocities in the specimen (and optionally wedge) are specified in /MFMC<m>/COMMON/ by the [2] attributes {SPECIMEN/WEDGE}\_VELOCITY. The first element is the shear velocity and the second is the longitudinal velocity.

#### Probe Placement (/MFMC<m>/PROBE\_\* and /MFMC<m>/COMMON/PROBE\_INDEX)

Each A-scan in each frame in a sequence corresponds to the same duration of ultrasonic time. This duration is assumed to negligible compared to the timescale associated with mechanical movement of the probe(s), hence the probe(s) is assumed to be at a stationary placement for each A-scan. Commonly, it will be the case that multiple A-scans (e.g. all those in a single frame) are recorded at the same probe placement. For this reason and to avoid unnecessary duplication, only unique probe placements are recorded in /MFMC<m>/ with the expanding [3,N\_Q<m>,N\_B<m>] datafields PROBE\_POSITION and PROBE\_{X/Y}\_DIRECTION. Each A-scan in each frame is cross-referenced into the probe-placement datafields through /MFMC<m>/PROBE\_PLACEMENT\_INDEX.

For all three probe-placement datafields the dimensions have the following interpretation. The first dimension, [3], refers to vector components in the GCS, and the third, expandable dimension, [N\_B<m>], is the probe placement index. The second dimension, [N\_Q<m>], is necessary because multiple probes with independent placements may be involved in a sequence (e.g. in a tandem configuration where one probe acts as a transmitter and one as a receiver). The size of this dimension is equal to the number of different probes used in the sequence and will often by unity. Indices in this dimension are cross referenced to the probe indices p in /PROBE<p> through /MFMC<m>/COMMON/PROBE\_INDEX.

The datafield, PROBE\_POSITION, provides the position vector, , of the origin of each PCS in the GCS for each probe placement index. The vectors, and , in PROBE\_{X/Y}\_DIRECTION specify the directions of the x- and y-axes of the PCS in the GCS. These should be orthogonal but if they are not, has priority and defines the x-axis direction of the PCS, while the component of that is orthogonal to defines the y-axis direction of the PCS. Note that the probe placement datafields work in a similar way to the ELEMENT\_POSITION and ELEMENT\_{MAJOR/MINOR} datafields in /PROBE<p>/, except that (a) they use the GCS rather than a PCS and (b) the magnitude of PROBE\_{X/Y}\_DIRECTION vectors does not have any physical meaning in contrast to the ELEMENT\_{MAJOR/MINOR} vectors.

\*probe placement

Fig. 3 Probe placement.

#### Other MFMC Sequence Datafields

\*todo for list below

/MFMC<m>/COMMON/TAG

/MFMC<m>/COMMON/DAC\_CURVE

/MFMC<m>/COMMON/RECEIVER\_AMPLIFIER\_GAIN

/MFMC<m>/COMMON/FILTER\_TYPE

/MFMC<m>/COMMON/FILTER\_PARAMETERS

/MFMC<m>/COMMON/FILTER\_DESCRIPTION

/MFMC<m>/COMMON/OPERATOR

/MFMC<m>/COMMON/DATE\_AND\_TIME

## Example Use Cases

### Single FMC frame with 1D linear array probe

In this example, the MFMC file contains the minimum information to record a single frame of FMC data acquired from a single linear array probe containing only 3 elements.

### Multiple FMC frames from 1D linear array probe

The set-up in this case is identical to that in the previous example and effectively this shows how multiple frames are added to a sequence.

### PWI frame

### \*that roof probe device

### \*pair of tandem probes with independent positions

\*need function in Matlab for dumping actual file contents and displaying. \*also write script for running these standard test cases and outputting results

# Appendix – Revision History