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# Direct sample positioning and alignment methodology for strain measurement by diffraction

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An ISO (International Organization for Standardization) TTA (Technology Trends Assessment) was published in 2001 for the determination of residual stress using neutron diffraction which identifies sample alignment and positioning as a key source of strain measurement error. Although the measurement uncertainty by neutron and synchrotron x-ray diffraction for an individual measurement of lattice strain is typically of the order of  $10-100\times10^{-6}$ , specimens commonly exhibit strain gradients of  $1000\times10^{-6}$  mm<sup>-1</sup> or more, making sample location a potentially considerable source of error. An integrated approach to sample alignment and positioning is described which incorporates standard base-plates and sample holders, instrument alignment procedures, accurate digitization using a coordinate measuring machine and automatic generation of instrument control scripts. The methodology that has been developed is illustrated by the measurement of the transverse residual strain field in a welded steel T-joint using neutrons. © 2005 American Institute of Physics. [DOI: 10.1063/1.1896726]

#### I. INTRODUCTION

The determination of the magnitude and distribution of residual stress in engineering components is important for many safety-critical applications. During manufacturing of these components, thermomechanical mismatches occur which lead to the existence of locked in or "residual" stresses. These stresses may be compressive or tensile. In general, tensile stresses are detrimental, supporting crack propagation for example, whereas compressive stresses are considered beneficial, limiting crack propagation. Because residual and applied stresses combine it is very important to know the residual stress field to accurately predict the total stress field and hence be able to determine component lifetimes and their creep behaviors.

The use of synchrotron x-ray and neutron facilities for the investigation of residual stresses in advanced engineering materials is increasing rapidly. This powerful technique essentially utilizes the crystal lattice of a polycrystalline material as a 3D strain gauge. Various types of neutron or x-ray sources are available worldwide. In the case of neutrons, sources are either reactor or accelerator (spallation) sources whereas intense high-energy x rays are provided by synchrotron sources.

An ISO (International Organization for Standardization) TTA (Technology Trends Assessment) was published in 2001 which describes standard protocols for the measurement of residual stress using neutrons<sup>2</sup> and work on an equivalent for the synchrotron x-ray technique has started. Standard procedures are necessary to ensure that residual stress measurements are reliable and repeatable. Standards are common in materials inspection and their adoption is important to establish academic and industrial confidence. One of the principal observations of these projects is the importance of accurate specimen location with respect to the measuring volume on the beamline. For an individual measurement of strain at a sample location, the specified uncertainty is often  $\approx \pm 50$  $\times 10^{-6}$ . This is typically the value obtained for a neutron measurement whereas  $\approx \pm 10 \times 10^{-6}$  is achieved for a synchrotron x-ray measurement. In a welded region<sup>3</sup> for example, the strain gradient may be as high as ≈1000  $\times 10^{-6}$  mm<sup>-1</sup> so that it is necessary to position the sample within  $\pm 50 \, \mu \text{m}$  to achieve the specified uncertainty. In the case of steep strain gradients the principal source of uncertainty is therefore often due to positioning errors. The need for efficient protocols for sample alignment and positioning was a key recommendation of the ISO/TTA. 2,4,5 Here we de-

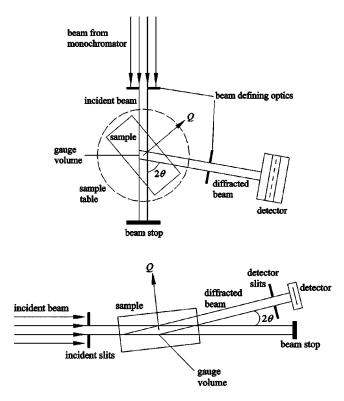


FIG. 1. Typical arrangement of neutron (top) and synchrotron x-ray (bottom) diffractometers for residual strain analysis.

scribe an integrated direct approach to sample alignment and positioning including a sample mounting system, instrument alignment procedure coupled with accurate sample digitization using a coordinate measuring machine (CMM) and automatic generation of instrument control script.

#### II. EXPERIMENTAL ISSUES

The design of neutron and synchrotron x rays diffractometers for strain measurement varies, although in most cases a "gauge volume," the volume from which strain is measured, is defined using beam optics on both the incident and diffracted beams. Figure 1 shows the outline diagrams of typical neutron and synchrotron x-ray diffractometers for strain analysis. The location, shape, and size of the gauge volume are defined by the diffraction angle and the beam optics system. In the case of neutron diffractometers, the angle between the incident and diffracted beam is usually around 90° thus obtaining a near-square shaped cross section. The gauge volume is elongated (diamond shaped) in the case of synchrotron x rays due to the high energy of the beam, which results in small Bragg angle values. The sample can be positioned with respect to the gauge volume using X, Y, Z translators, and a rotation table is used to adjust the strain measurement direction.

For most instruments an area of uncertainty is the interface between the diffractometer and the sample. Often there is not a standard or adaptable mounting system that enables reproducible sample positioning on the instrument. Users often have to adopt a temporary system such as sticky tape or glue. These arrangements increase positioning uncertainty, irreproducibility in sample positioning, and inconvenient and

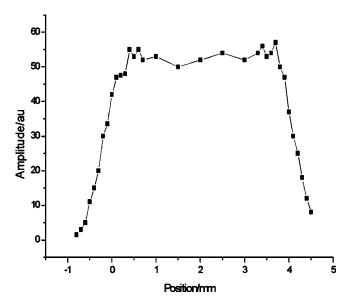


FIG. 2. Entrance scan showing the intensity profile as a sample is scanned across the gauge volume.

substantial time delays. Various solutions have been proposed to overcome these issues at neutron and synchrotron x-ray facilities. These generally involve a reliable sample fixturing system such as a large accurate breadboard at the sample position,<sup>6</sup> kinematic sample mounts<sup>7</sup> or the use of a CMM for sample metrology.<sup>8</sup> A further problem is that although the gauge volume is defined using beam optics, its position relative to the reference point of the diffractometer has often not been established accurately. In these circumstances, a sample surface is located through the use of "entrance and exit scans" in which the sample is traversed through the gauge volume and the diffracted intensity is recorded. Typical entrance and exit scans are shown in Fig. 2 for a relatively thin sample ( $\approx 4$  mm thick). If the gauge is fully within the sample, the diffracted intensity is constant, neglecting attenuation effects due to different path lengths. The edge of a sample is defined as the point when half of the gauge volume is immersed in the sample therefore giving half the maximum diffracted intensity. This process is time consuming, particularly using neutrons, and is of limited assistance when the sample geometry is complex, when a sample has to be rotated or when the count rates are low. Furthermore, texture variations and strain changes can result in misleading scattered intensity entrance (or exit) profiles.

In order to overcome these issues a "direct positioning" approach has been adopted which provides a significant improvement in accuracy as well as minimizes the time needed for sample alignment and therefore improves experimental beamtime efficiency. The direct positioning concept is based around the provision of a standard, accurate and reliable physical interface between the sample and beamline, together with accurate determination of sample geometries using a coordinate measuring machine, determination of the spatial relationship between sample and gauge volume and the implementation of automated script generation for sample stage movement on the diffractometer.

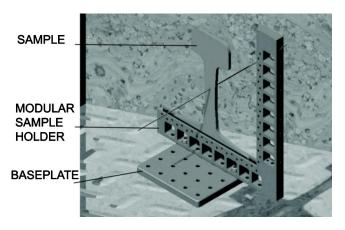


FIG. 3. View of a rail specimen mounted in a sample holder and attached to a base-plate.

#### **III. SAMPLE POSITIONING SYSTEM**

Sample fixturing benches are common in a range of scientific metrology applications. These are often based on a threaded or dowelled breadboard such as optical benches. In order to implement a direct positioning system, an accurate sample positioning system was designed and tested. The sample is mounted in a sample holder that matches a baseplate. The base-plates were designed to replace, or attach to, the top plate of the diffractometer. The base-plate system consists of 12 mm thick aluminium alloy plates with a matrix of 25 mm pitch holes. Each hole is half threaded (M6 threaded holes on a square matrix at  $25\pm0.005$  mm centers) and half reamed (H5) to take a 6 mm (m6) dowel for precise positioning, or the straight shank of an M6 screw for routine fixing.

Sample holders with a matching 6 mm hole pattern have been designed to hold flat samples (such as slices of railway sections, welds,...), complex-shaped samples (turbine blades,...) and pin-shaped specimens which allow for gauge volume position definition. A modular set of sample holders was designed in which two similar units may be combined to form an "L-shaped" sample holder. The interest of this design is that it enables the user to accurately rotate the sample through 90° steps without dismounting it. This rotation is useful for stress determination. Figure 3 shows a view of a rail section mounted in an "L-shaped" sample holder. The holder is shown mounted upon a small base-plate. These sample holders are compatible with the base plates and enable accurate and repeatable positioning of the sample on the base plate. It enables samples to be rapidly positioned to within  $\pm 100$  microns and oriented to within  $\pm 0.1^{\circ}$ , and to within  $\pm 10$  microns and  $\pm 0.02^{\circ}$  with care. Diffractometers at the ILL-ESRF (Grenoble, France) and the coordinate measuring machine at the FaME38 lab (Grenoble, France) have been equipped with this positioning system.

## IV. BEAMLINE PREPARATION AND SAMPLE DIGITISATION

Coordinate measuring machines (CMM) are routinely used in industry as quality control equipment (inspection) or less commonly used for model creation (digitization). After mounting on a standard base plate, samples are digitized

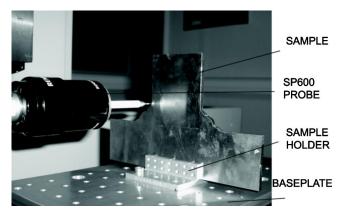


FIG. 4. A weld section under investigation using the SP600 scanning probe. The sample is mounted in a standard sample holder and a base-plate is available on the CMM base.

with a CMM prior to an experiment. The FaME38 laboratory is equipped with a Mitutoyo EURO-C-A776 CMM. A Renishaw TP20 or SP600 ruby tipped probe is mounted on a motorized Renishaw PH10MQ articulated head that can be orientated in any direction over a hemisphere (Fig. 4). The TP20 probe is designed for discrete measurements whereas the advanced SP600 probe is able to *scan* a surface whilst maintaining contact with it with an accuracy of  $\approx \pm 2~\mu m$ . A Nextec Wizprobe laser measuring system is also available for non-contact measurements.

The direct positioning concept minimizes the amount of experimental time used for sample alignment and positioning, as the majority of procedures for sample setup (alignment and positioning) are performed off-line before an experiment. At the start of an experimental session a pin-shaped specimen is placed in a standard pin holder. The direct positioning procedure starts by setting the top-center of the pin specimen as the origin of the measurements on the CMM. The orthogonal coordinate system on the CMM is chosen so as to be inverse of the *X*, *Y*, *Z* translator system on the beamline. In effect, the axes on the CMM are the same as those used on the instrument but in the *opposite* direction.

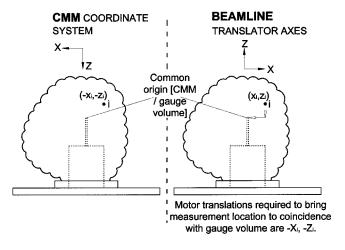


FIG. 5. Simplified 2-axis representation of the transposition of CMM coordinates into motor translations on the beamline. Through the adoption of an inverse coordinate system, a sample location (-Xi,-Zi) can be located at the gauge volume through a beamline motor translation of -Xi,-Zi. The position of the pin and pin holder is shown by dashed lines.

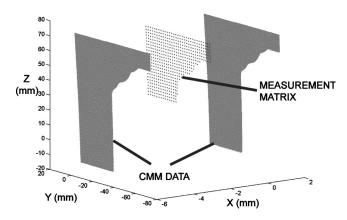


FIG. 6. Measurement points generated automatically using CMM surface data.

Standard base plates are used as a reference to align the coordinate systems on the diffractometer and on the CMM. The top centre of the pin is used as a *common reference point* on the beamline (coincident with the center of the gauge) and on the CMM (origin of the measurements). The sample to be studied is mounted in a standard sample holder and is then digitized in the coordinate system on the CMM.

The next stage of the procedure is to position the gauge volume at the top centre of the pin. This can be achieved in different ways depending on the design of the instrument. For the neutron case, the pin is often placed at the centre of rotation of the goniometer and the gauge volume is made coincident with the tip of the pin by translating the primary and secondary optics across the incident and diffracted beam respectively. Variations in the intensity of the scattered beam are used to locate the beam with respect to the pin. For the synchrotron x-ray case, the pin is translated across the beam and the variations in scattered intensity enable the tip of the pin to be positioned at the gauge volume. In both cases, when the tip of the pin and the gauge volume are coincident, the translation axes X, Y, and Z of the diffractometer are set to zero.

At this stage the gauge volume and the top center of the pin are coincident with the reference point on the beamline. Coordinates obtained from a real sample using the CMM can now be used to drive the X, Y, Z translators and position the desired sample location at the center of the gauge volume. During the sample transfer from the CMM to the beamline, the relative position of the sample holder with respect to the pin holder position on the base plate (and the sample within the sample holder) must remain fixed or the coordinates are not correct. Figure 5 shows a simplified 2D drawing of the transposition of CMM coordinates into motor translations on the beamline. If we consider one required measurement point i in the sample, the coordinates of this point in the CMM coordinate system  $(X_{CMM} = -X_i, Z_{CMM} = -Z_i)$  and beamline coordinate system  $(X_{\text{Beamline}} = X_i, Z_{\text{Beamline}} = Z_i)$  are opposite. The motor translations required on-line to make this point coincident with the centre of gravity of the gauge volume  $(X_{\text{Beamline0}}=0, Z_{\text{Beamline0}}=0)$  correspond to translator displacements of  $(X_{\text{Beamline}} - X_{\text{Beamline}} = -X_i, Z_{\text{Beamline}} - Z_{\text{Beamline}})$  $=-Z_i$ ). It must be remembered that in order to locate the gauge volume at a sample location it is necessary to move

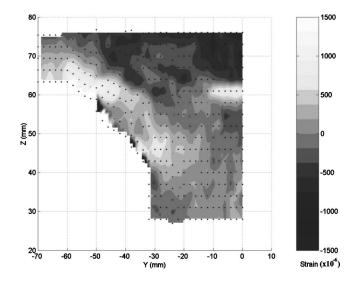


FIG. 7. Transverse residual strain data for steel T-welded sample.

the *sample* and not the *gauge volume* which is fixed in space at the reference point. Therefore, the coordinates obtained from sample digitization on CMM can directly be used as the input motor positions to place a measurement point in the sample at the centre of the gauge volume.

Most of the stages of this procedure are performed offline. The only preliminary neutron measurement needed is the positioning of the gauge at the top centre of the pin, which is done just once at the start of an experimental session. Digitized samples in standard holders can then be directly aligned and positioned with respect to the reference point on the beamline.

#### V. AUTOMATED MACRO GENERATION

Experiments to measure residual strain profiles usually require many measurements over an area or volume. For large numbers of measurement points it is unreliable and time-consuming to input by hand the commands to move the translators. By manipulating the coordinates obtained by digitization of the sample via software, it is possible to automatically write the script (series of commands sent to the translation and rotation axes and to the detectors) to carry out the neutron or synchrotron x-ray strain mapping within the volume of the sample. The time spent on the beamline to set up the strain mapping experiment is then reduced to a minimum and the possibilities of errors are reduced.

A program named *Easyscript* was developed for beamlines at the ILL and ESRF using the Matlab computing language. CMM data are used to generate the series of points where neutron or synchrotron x-ray measurements are to be performed in the bulk of the sample according to the requirements of the user. A script for the experiment is then generated. This includes the motor movements needed to place the desired location in the sample at the gauge volume and the acquisition commands for the detector. The software takes in account the translation limits of the motors and removes the points that are outside these machine limits. An option is available which prevents measurements which straddle a surface to prevent possible surface aberrations.<sup>11</sup>

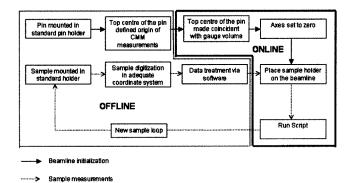


FIG. 8. Principal operations of the direct sample positioning procedure.

#### VI. APPLICATION OF DIRECT SAMPLE POSITIONING

The protocols and software were tested using a fusion welded T-joint made of ferritic steel (see Fig. 4). The experiment was performed using the strain diffractometer D1A at the ILL neutron source in Grenoble, France. The welded section was digitized on both sides using the CMM with a high resolution measurement pitch  $(0.2 \times 0.2 \text{ mm}^2)$ . An iron pin was used to define the reference point on the beamline. *Easyscript* was then used to generate a square grid of points at mid-thickness in the sample where neutron measurements were to be undertaken. Figure 6 shows the resulting map where neutron measurements were taken. The coordinates of these points were used to automatically write the series of commands that were sent to the motors/detector controlling units during the strain mapping experiment.

The sample was then mounted on the diffractometer table using the standard positioning system. Data analysis was performed using the software package LAMP developed by the ILL computing service and the strain results of the experiment are shown in Fig. 7.

#### VII. DISCUSSION

The need for improved sample positioning procedures for residual strain measurements has been identified in the ISO TTA (2001). The methodology developed uses a rigorous alignment procedure and incorporates the novel use of a CMM for accurate sample digitization. Figure 8 shows the principal operations of the developed procedure. It is possible to perform the majority of operations offline, before the start of an experiment. Indeed, if required, users are able to measure all samples using the CMM and generate the control scripts in advance of the beamline initialization. The standard base-plates and sample holders give reliable fixturing and allow users to easily create their own mounting system prior to an experiment. The Easyscript software package facilitates automatic script writing which reduces the probability of errors and saves time. Measurements are made only at defined positions within a sample, whereas in the past it has often been common practice to define a rectangular measurement matrix that extends past the limits of a complex shaped sample. These developments are now available for residual strain experiments at the ILL and ESRF beamlines in Grenoble, France. This direct positioning concept using a coordinate measuring machine for sample digitization and alignment is likely to be adopted for residual strain measurement at international neutron and synchrotron x-ray sources. However the methodology is more widely applicable to any neutron, synchrotron x-ray or lab-based source where the accurate location of specimens is an important consideration.

Although the source of many errors has been reduced, there are further issues which are currently being addressed. The space around a sample is often limited; this is particularly the case with neutron measurements as the beam defining slits are placed close to the sample. It is important to eliminate sample collisions with the beam defining optics as this often necessitates a complete realignment. The concept that the experiment should be simulated offline is currently being pursued. By using the CMM we can essentially obtain a CAD knowledge of the sample and by combining this with a CAD model of the beamline and the desired measurement script, it is possible to create a *virtual beamline*. In this way it is possible to predict and prevent collisions. This type of simulation will further enhance the experimental efficiency of residual strain experiments.

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