# A *p*-ADIC RANSAC ALGORITHM FOR STEREO VISION USING HENSEL LIFTING

#### PATRICK ERIK BRADLEY

ABSTRACT. A *p*-adic variation of the Ran(dom) Sa(mple) C(onsensus) method for solving the relative pose problem in stereo vision is developped. From two 2-adically encoded images a random sample of five pairs of corresponding points is taken, and the equations for the essential matrix are solved by lifting solutions modulo 2 to the 2-adic integers. A recently devised *p*-adic hierarchical classification algorithm imitiating the known LBG quantisation method classifies the solutions for all the samples after having determined the number of clusters using the known intra-inter validity of clusterings. In the successful case, a cluster ranking will determine the cluster containg a 2-adic approximation to the "true" solution of the problem.

Key words: *p*-adic classification; relative pose; essential matrix; RANSAC

## 1. INTRODUCTION

High dimensional and sparse encodings of data tend to be ultrametric, and ultrametric spaces allow certain computational operations, like nearest neighbour finding, to be carried out very efficiently [17]. This suggests, for the task of hierarchical classification, an ultrametric encoding of data from the start. This has the advantage that the hierarchical structure is uniquely determined by the ultrametric, however at the price of having to find a suitable encoding in an ultrametric space [4]. A natural family of ultrametric spaces is given by  $\mathbb{Q}_p$ , the *p*-adic numbers for any prime number *p* of choice. A first application of classification algorithms to *p*-adic data in image segmentation is described in [1], where it was found that the *p*-adic algorithms outperformed their classical counterparts in efficiency. In [5], it was observed that *p*-adic clustering algorithms need not change the metric when computing distances between clusters.

The task of finding optimal classifications lead to the well-known LBG algorithm, named after the initials of their authors [15]. The clusters from the previous step are split by regrouping the data around the new "centres" in an optimal way. A direct *p*-adic analogue does not exist. However, if clusters are interpreted as vertices in the dendrogram for the given data, then splitting can be performed by replacing a vertex by its children. Splitting in direction of highest gain and, after finding the clustering, determining cluster centres leads to what we call LBG<sub>*p*</sub> algorithm [3]. For LBG, the desired number of clusters has to be pre-specified. For LBG<sub>*p*</sub>, we have a pre-specified upper bound for the cluster number. This leads to the issue of determining that number or bound. In [20], an optimisation scheme for the number of clusters is developped for the *k*-means clustering algorithm. We propose a *p*-adic

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adaptation of this for choosing optimal clusterings among the  $\text{LBG}_p$ -optimal ones of varying size.

An important algorithm in image analysis is Random Sample Consensus, known as RANSAC [7]. In short, this is a general method for fitting a model to experimental data by randomly sampling the minimal number of points necessary for the fit. Then that set of feasible points is enlarged by adding all nearby points, i.e. those points having a fitted model not much different from the first fit. This is the consensus set. The sample with largest consensus set yields the best model prediction. A variation of this idea would be to perform a classification of the fitted models for all random samples taken from the data. Then the biggest cluster (with respect to some measure) contains in its centre the "true" model. Here, we describe a *p*-adic form of this random sample consensus via classification using the LBG<sub>p</sub> algorithm, applied to the relative pose problem in stereo vision as described in the following paragraph.

The issue of estimating camera motion from two views is classical by now, and methods from projective and algebraic geometry towards its solution were employed already at an early stage. The relationship between the views is established by finding correspondences between point pairs taken from both images. The fundamental matrix faithfully encodes the geometric relationship between the two images. For normalised cameras, the fundamental matrix coincides with the so-called essential matrix. In general, the two matrices are related through the camera calibration. Hence, if the calibration is known, it is sufficient to estimate the essential matrix in order to solve the relative pose problem up to a sign ambiguity. In order to efficiently use a RANSAC for this task, it makes sense to use in each sample the minimal number of point correspondences needed. This number is known to be five [13]. Only recently, an efficient solution to the five-point relative pose problem has been developped [18]. The first algorithms use eight point pairs, for which all equations are linear [16]. Seven point pairs require a non-linear constraint, and so does the method using six points [9, 11, 19]. The idea for the five point problem is to first solve the five linear constraints, and then insert the general solution into the other constraints. This leads to nine homogeneous cubic equations in four unknowns. These describe a subspace S of projective three-space  $\mathbb{P}^3$  over the ground field K, and the relative pose problem has a solution if the space S is zero-dimensional. In that case, it is known to be of cardinality ten [6], if multiplicities are taken into account and all solutions from an algebraic closure of K are allowed. Nistér's approach is to eliminate variables and then numerically solve the resulting univariate polynomial of degree 10. An alternative natural approach is by using Gröbner bases [21]. A formulation of the problem as a polynomial eigenvalue problem which can be solved robustly and efficiently is found in [14].

By considering image coordinates as 2-adic numbers, e.g. through an interval subdivision process, we can rewrite the equations for the five-point relative pose problem as polynomials with coefficients in  $\mathbb{Z}_2$ , the 2-adic integers. This allows us to use Hensel's lifting method for their solution, and we arrive at 2-adic essential matrices which can be approximated by finite 2-adic expansions depending on the desired resolution. In this article, we take a closer look at the structure of the cubic equations and arrive at a union of algebraic varieties defined by linear and quadratic equations which have to be intersected with the remaining cubic constraint over the field  $\mathbb{F}_2$  in the initial stage before the lifting. An efficient solution of those particular

equations with Gröbner bases is deferred to future work. However, we expect the existing literature on Gröbner basis methods for equations over finite fields to provide results which can be "tuned" to our situation. The advantage of using the field  $\mathbb{F}_2$  is that Gröbner basis computations become very efficient. Together with our decomposition into quadratic and linear equations, we expect higher performance in comparison with the real methods operating on the undecomposed equations. Using the Hensel lifting methods leads us to expect higher robustness. However, these expectations yet await practical evaluation.

The following section collects general facts on p-adic numbers which we will use, and fixes some notation. For the convenience of the reader, we prove that p-adic vectors can be viewed as p-adic numbers in an unramified extension field of  $\mathbb{Q}_p$  via an isometric isomorphism. In Section 3, we review the LBG<sub>p</sub> algorithm and show how one can determine the number of clusters. Section 4 explains how to arrive at a 2-adic encoding of image coordinates, and then develops the lifting algorithm for the five-point relative pose equations. Section 5 incorporates that algorithm into RanSaC<sub>p</sub>, the random sample consensus algorithm via classification.

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### 2. Generalities

In this section, we review some facts about p-adic numbers for later use. A general introduction can be found in [8], the Haar measure on local fields is introduced in [22].

2.1. *p*-adic fields. Let *p* be a prime number, and *K* a field which is a finite extension field of the field  $\mathbb{Q}_p$  of rational *p*-adic numbers. We call *K* a *p*-adic field, and its elements simply *p*-adic numbers. *K* is a normed field whose norm  $| |_K$  extends the *p*-adic norm  $| |_p$  on  $\mathbb{Q}_p$ . Let  $\mathcal{O}_K := \{x \in K \mid |x|_K \leq 1\}$  denote the local ring of integers of *K*. Its maximal ideal  $\mathfrak{m}_K = \{x \in K \mid |x|_K < 1\}$  is generated by a *uniformiser*  $\pi$ . It has the property  $v(\pi) = \frac{1}{e}$ , where  $e \in \mathbb{N}$  is the ramification degree of  $K/\mathbb{Q}_p$ . If e = 1, then *K* is called *unramified* over  $\mathbb{Q}_p$ .

All elements  $x \in K$  have a  $\pi$ -adic expansion

(1) 
$$x = \sum_{i \ge -m} \alpha_i \pi^i$$

with coefficients  $\alpha_i$  in some set  $\mathcal{R} \subseteq K$  of representatives for the residue field  $O_K/\mathfrak{m}_K \cong \mathbb{F}_{p^f}$ . In the case f = 1, the choice  $\mathcal{R} = \{0, 1, \ldots, p-1\}$  is quite often made. If K is unramified of degree n over  $\mathbb{Q}_p$ , then f = n.

The Haar measure on K will be denoted by dx and is normalised to

$$\int_{O_K} dx = 1.$$

2.2. *p*-adic vectors as *p*-adic algebraic numbers. In this subsection, we show how to consider higher-dimensional *p*-adic data as one-dimensional data in some appropriate unramified field extension. As a consequence, any classification method with *p*-adic numbers can be applied to *p*-adic vector data.

On the vector space  $\mathbb{Q}_p^n$  there is the maximum norm  $\|\cdot\|_{\max}$  given by

$$||x||_{\max} = \max\left\{|x_1|_p, \dots, |x_n|_p\right\},\$$

where  $x = (x_1, \ldots, x_n) \in \mathbb{Q}_p^n$ . The following lemma allows to consider vectors as one-dimensional objects in some *p*-adic number field:

**Lemma 2.1.** There is an isometric isomorphism between normed vector spaces:

$$(\mathbb{Q}_p^n, \left\|\cdot\right\|_{\max}) \cong (K, \left|\cdot\right|_K),$$

where K is any unramified extension field of  $\mathbb{Q}_p$  of degree n. Furthermore, for all n there exists such a p-adic field K.

*Proof. Isometry.* Any  $x \in K$  has a *p*-adic expansion

(2) 
$$x = \sum_{\nu \ge m} a_{\nu} p^{\nu},$$

where p retains the uniformising property, because K is unramified over  $\mathbb{Q}_p$ . The coefficients  $a_{\nu} \in K$  are taken from a set  $\mathcal{R}$  of representatives of the residue field  $\kappa = O_K/pO_K \cong \mathbb{F}_{p^n}$ , where  $\mathcal{R}$  contains the zero element of K. The residue field, as a vector space, is isomorphic to  $\mathbb{F}_p^n$ . Hence, the coefficients can be identified with vectors in  $\mathbb{Q}_p$  whose entries are taken from the set  $\{0, \ldots, p-1\}$  which represents the residue field  $\mathbb{F}_p$ . This yields a bijection between K and  $\mathbb{Q}_p^n$  which is in fact an isomorphism of vector spaces. Observe now that the norm  $|x|_K = p^{-m}$ , where  $m \in \mathbb{Z}$  is the smallest exponent  $\nu$  of p in the p-adic expansion (2) of x such that  $a_{\nu} \neq 0$ . By the above identification of x with a vector  $(x_1, \ldots, x_n) \in \mathbb{Q}_p^n$ , this means that m is the smallest exponent for which the coefficient is not the zero vector. Hence,

$$|x|_{K} = p^{-m} = ||(x_{1}, \dots, x_{n})||_{\max}$$

as asserted.

*Existence.* Let  $\zeta$  be a primitive  $(p^n - 1)$ -th root of unity. Then the cyclotomic field  $K = \mathbb{Q}_p(\zeta)$  is an unramified extension field of  $\mathbb{Q}_p$  of degree n [8, Prop. 5.4.11].  $\Box$ 

Consequently, a set of *n*-dimensional *p*-adic vectors can be treated as data taken from a one-dimensional algebraic *p*-adic field K. Hence, its dendrogram for the maximum norm can be viewed as a subtree of the Bruhat-Tits tree for K. In particular, the classification methods from [3, 5] for data from K apply to this case.

### 3. An Almost optimal *p*-adic classification algorithm

After briefly reviewing the *p*-adic variation of the hierarchical classification algorithm of [15], we show how to determine the optimal number of clusters in the *p*-adic case. More details on the *p*-adic classification algorithm can be found in [3]. The cluster number determination is a *p*-adic analogue of the method from [20].

3.1.  $\mathbf{LBG}_p$ . In [15] a clustering algorithm is presented which determines optimal clusterings of given real vector data. This so-called *split-LBG algorithm* constructs cluster centres around which the data are grouped in an optimal manner. This method has no direct *p*-adic analogon. However, [3] shows an adaptation of the split-LBG algorithm to *p*-adic data which locally minimises the expression

$$E(\mathscr{C}, \mathbf{a}) = \sum_{c \in \mathscr{C}} \sum_{x \in C} |x - a_C|_K,$$

where  $\mathscr{C}$  is a partition (clustering) of given data X, and  $\mathbf{a} = (a_C)_{C \in \mathscr{C}}$  consists of centres of clusters C. The clusterings are bounded a priori in size by  $|\mathscr{C}| \leq k$ . The method is to first subdivide given clusters in order to obtain largest decrease in  $E(\mathscr{C}, \mathbf{a})$ , and then within the found clustering to find centres in a second step. The centres are characterised by their minimising property for  $E(\mathscr{C}, \mathbf{a})$  with a given clustering  $\mathscr{C}$ . More details can be found in [3]. The cluster centres will be used later in Section 5 for obtaining an estimated solution to the relative pose problem in stereo vision.

3.2. Determining the number of clusters. The  $\text{LBG}_p$  algorithm as described in the previous subsection uses as input a pre-specified upper bound for the number of clusters. Usually, however, this number is not known a priori. There are in the literature various methods for finding optimal cluster numbers. The application we have in mind is to find a clustering in which one cluster contains the "best" approximations to some unknown quantity, and the other clusters are "outliers". Hence, the ideal clustering should contain compact clusters which are well separated. For this reason, we define a *p*-adic version of the intra-inter-validity index from [20].

Let  $X \subseteq K$  be a finite set, and  $\mathfrak{X}_k(X)$  the set of all verticial clusterings of X with cardinality  $\ell \leq k$ . For a choice of cluster centres  $\mathbf{a} = (a_C)_{C \in \mathscr{C}}$  in a given clustering  $\mathscr{C} \in \mathfrak{X}_k(X)$ , we define the following quantities:

$$\begin{aligned} \operatorname{Intra}(\mathscr{C}) &= \frac{1}{|X|} \sum_{C \in \mathscr{C}} \sum_{x \in C} |x - a_C|_K \\ \operatorname{Inter}(\mathscr{C}) &= \min_{C \neq C' \in \mathscr{C}} |a_C - a_{C'}|_K \\ \operatorname{Validity}(\mathscr{C}) &= \frac{\operatorname{Intra}}{\operatorname{Inter}} \end{aligned}$$

Lemma 3.1. Intra and Inter do not depend on the choice of cluster centres a.

*Proof.* Intra. This follows from the definition of cluster centre, as

Intra
$$(\mathscr{C}) = \frac{1}{|X|} E(\mathscr{C}, \mathbf{a}).$$

Inter. This follows from the strict triangle inequality, as distinct clusters  $C \neq C'$  from  $\mathscr{C}$  are disjoint.

The function Intra measures the compactness of a cluster, whereas Inter is a measure for the inter-cluster distance. A "good" clustering would minimise Intra and maximise Inter. So, an obvious measure combining both tasks is given by Validity as the ratio of both measures. **Definition 3.2.** The quantity

$$\operatorname{vi}_k(X) = \min_{\mathscr{C} \in \mathfrak{X}_k(X)} \operatorname{Validity}(\mathscr{C})$$

is called the k-th validity index of the data X.

**Lemma 3.3.** If N = |X|, then  $vi_N(X) = 0$ .

*Proof.* This follows from the fact that  $\mathfrak{X}_N$  contains the clustering consisting of N singletons.

Hence, the ideal cluster number is to be determined by computing the k-th validity index for  $k \ll N$ .

**Remark 3.4.** It is to be expected that for genuine data X, the decreasing function  $k \mapsto \operatorname{vi}_k(X)$  remains constant on some large interval I inside  $\{2, \ldots, |X|\}$ , and that the minimum of Validity on  $\mathfrak{X}_k$  with  $k \in I$  is attained on some clustering  $\mathscr{C} \in \mathfrak{X}_k$  such that  $|\mathscr{C}| \ll |X|$ .

**Definition 3.5.** A clustering C of minimal validity in the sense of Remark 3.4 is called an ideal clustering.

## 4. The dyadic 5-point relative pose equations from stereo vision

A first *p*-adic formulation of the *n*-point relative pose problem of stereo vision is presented in [2]. There, the equations are derived from a *p*-adic projective camera model. Here, we first review the 2-adic image encoding from a hierarchical classification point of view, and then propose a refined lifting algorithm from the reduced equations modulo 2 to  $\mathbb{Z}_2$ -rational solutions.

4.1. 2-adic image encoding. Let  $R \subseteq \mathbb{R}^n$  be the unit hypercube  $[0,1]^n$ . It can be subdivided by n hyperplanes parallel to the coordinate hyperplanes into  $2^n$ hypercubes of equal volume. Each of these smaller hypercubes can be subdivided into even smaller hypercubes in the same way. Repeating this process yields an infinite rooted tree  $\mathscr{T}$  whose vertices are those hypercubes, and edges are given by pairs  $(R_n, R_{n+1})$  of hypercubes where  $R_{n+1}$  is one of the parts of  $R_n$  obtained in the *n*-th subdivision step. The root of  $\mathscr{T}$  is given by  $R_0 = R$ , and each vertex has  $2^n$  children vertices.

Let v be a vertex of  $\mathscr{T}$ , and denote ch(v) the set of its children vertices. A family  $\chi$  of bijections

$$\chi_v \colon \operatorname{ch}(v) \to \{0,1\}^n$$

defines a labelling  $\lambda_{\chi}$  on Edges( $\mathscr{T}$ ), the set of edges of  $\mathscr{T}$ :

$$\lambda_{\chi} \colon \mathrm{Edges}(\mathscr{T}) \to \{0,1\}^n, \quad (v,w) \mapsto \chi_v(w),$$

and this allows for a 2-adic encoding of R, as will be seen in the following.

First, observe that an end of  $\mathscr{T}$ , i.e. an infinite (injective) path beginning in  $R_0$ , corresponds to a decreasing sequence of hypercubes

$$R_0 \supseteq R_1 \supseteq R_2 \supseteq \ldots$$

having a limit

$$\bigcap_{\nu \in \mathbb{N}} R_{\nu} = \{r\}$$

with a well defined point  $r \in R$ . This yields an inclusion map

$$\iota: \operatorname{Ends}(\mathscr{T}) \to R,$$

where  $\operatorname{Ends}(\mathscr{T})$  is the set of ends of  $\mathscr{T}$ .

We can view an end  $\gamma$  as a tree whose edges  $e_0, e_1, e_2, \ldots$  are directed away from the root  $R_0$ . They can be numbered by saying that  $\nu(e)$  is the number of edges on the path segment  $[R_0, o(e)]$ , where o(e) is the origin vertex of e. Now, traversing down a path  $\gamma \in \text{Ends}(\mathscr{T})$  and picking up the labels on edges  $e \in \text{Edges}(\gamma)$  along the path yields a 2-adic vector, as given by the map

$$\varpi_{\chi} \colon \operatorname{Ends}(\mathscr{T}) \to \mathbb{Z}_2^n, \quad \gamma \mapsto \sum_{e \in \operatorname{Edges}(\gamma)} \lambda_{\chi}(e) \, 2^{\nu(e)},$$

and which can be interpreted as an algebraic p-adic number in some unramified p-adic field K by Lemma 2.1.

**Lemma 4.1.** The map  $\varpi_{\chi}$  is bijective, and there exists a labelling  $\lambda_{\chi}$  such that  $\iota \circ \varpi_{\chi}^{-1}$  coincides with the map

$$\mu_2 \colon \mathbb{Z}_2^n \to R, \quad a = \sum_{\nu \in \mathbb{N}} \alpha_\nu \, 2^\nu \mapsto \sum_{\nu \in \mathbb{N}} \alpha_\nu 2^{-(\nu+1)},$$

with  $a_{\nu} \in \{0, 1\}^n$ .

*Proof.*  $\varpi_{\chi}$  bijective. A 2-adic vector in  $\mathbb{Z}_2^n$  corresponds uniquely to a sequence  $\lambda_0, \lambda_1, \lambda_2, \ldots$  of elements from  $\{0, 1\}^n$ . This in turn corresponds uniquely to a path from  $R_0$  by construction of the labels on  $\operatorname{Edges}(\mathscr{T})$ . Hence,  $\varpi_{\chi}$  is bijective.  $\mu_2$ . Let  $\chi^{\mu}$  be the family of bijections

$$\chi^{\mu}_{v} : \operatorname{ch}(v) \to \{0,1\}^{n}$$

given by the following construction. Consider the case n = 1. Then  $\mathscr{T}$  is a binary tree in which for given vertex v any  $w \in ch(v)$  corresponds to the interval  $R_w$ which is either the left or the right half of the interval  $R_v \subseteq [0, 1]$  corresponding to v. Now, by defining

$$\chi_v^{\mu} \colon \operatorname{ch}(v) \to \{0, 1\}, \quad w \mapsto \begin{cases} 0, & R_w \text{ is the left half of } R_v \\ 1, & R_w \text{ is the right half of } R_v \end{cases}$$

we obtain the labelling  $\lambda_{\mu} := \lambda_{\chi^{\mu}}$ , and the map  $\varpi_{\mu} := \varpi_{\chi^{\mu}}$ . We need to prove that

$$\bigcap_{v \in \operatorname{Vert}(\gamma)} R_v = \{\mu_2(\varpi_\mu(\gamma))\}$$

for all  $\gamma \in \text{Ends}(\mathscr{T})$ . Let  $\gamma_{\nu}$  be the segment of  $\gamma$  given by the first edges  $e_0, \ldots, e_{\nu}$ . The terminal vertex of  $\gamma_{\nu}$  corresponds to an interval  $R_{\nu+1}$  of length  $\frac{1}{2^{\nu+1}}$ . Let  $x_{\nu+1}$  be the left boundary of  $R_{\nu+1}$ . Inductively, it can be seen as given by

$$x_{\nu+1} = x_{\nu} + \epsilon_{\nu} \cdot \frac{1}{2^{\nu+1}},$$

where  $x_{\nu}$  is the left boundary of  $R_{\nu}$  and  $\epsilon_{\nu} \in \{0, 1\}$ . Clearly, it holds true that

$$\epsilon_{\nu} = \chi^{\mu}_{v_{\nu}}(v_{\nu+1}) = \lambda_{\mu}(e_{\nu}),$$

if edge  $e_{\nu}$  is given as  $e_{\nu} = (v_{\nu}, v_{\nu+1})$ . Hence,

$$\varpi_{\mu}(\gamma) = \sum_{\nu \in \mathbb{N}} \epsilon_{\nu} 2^{\nu}$$

and it follows that the sequence  $(x_{\nu})$  converges with respect to the Euclidean absolute norm to

$$\sum_{\nu \in \mathbb{N}} \epsilon_{\nu} 2^{-(\nu+1)} = \mu_2(\varpi_{\mu}(\gamma)),$$

as asserted.

The general case follows from the case n = 1 by applying it to each individual coordinate.

Since  $\mu_2$  is a bijection, it follows that  $\iota$  is bijective, too.

**Remark 4.2.** A rectangular photographic image can be viewed as a rectangular domain in  $\mathbb{R}^2$ . A digital image, however, will be represented for simplicity by an  $N \times N$  grid in  $\mathbb{R}^2$  in which we may assume that  $N = 2^n$ . Hence, the points on the image grid correspond bijectively to the elements of  $\mathbb{Z}/2^n\mathbb{Z} \times \mathbb{Z}/2^n\mathbb{Z}$ , and the exponent n defines the resolution of the image. The subdivision process as before increases each coordinate resolution by 1, and only the physical restrictions prevent us from obtaining a 2-adic image grid  $\mathbb{Z}_2 \times \mathbb{Z}_2$ . Hence, we may assume two idealised encodings of the digital image square: the Archimedean one is a real square given by the unit square  $[0,1]^2 \subseteq \mathbb{R}^2$ , and the p-adic encoding is  $\mathbb{Z}_2 \times \mathbb{Z}_2$ . The two ideal encodings are assumed compatible in the sense that the Monna map  $\mu_2 \colon \mathbb{Z}_2^2 \to \mathbb{R}^2$ embeds the one into the other. Physically, the real or 2-adic coordinates will be approximated in finite resolution by a grid isomorphic to  $\mathbb{Z}/2^n\mathbb{Z}$  with varying (and ideally arbitrary) n.

4.2. The linear and cubic equations. Assume that there are two views on a static 3D scene taken by cameras with known calibration. Here, the camera model is projective, and we briefly explain how the equations for estimating the geometric relationship between the two 2D views are derived. For an introduction to multiple view geometry we refer to [10].

A projective camera is a projective map  $\mathbb{P}^3 \to \mathbb{P}^2$ . If two such maps are given, the geometric relationship between the two images I and I' allow to estimate a recontsruction of the 3D scene. This relationship is given by the so-called *essential matrix* E, a projective  $3 \times 3$  matrix satisfying

(3) 
$$u_i^T E u_i' = 0$$
  $(i = 1, ..., N),$ 

where  $u_i \in I$  and  $u'_i \in I'$  are image points corresponding to the same point in 3-space. They are given as vectors with 3 homogeneous coordinates. The equations (3) are linear in the 9 unknown entries of E. Reconstruction from E is possible through the factorisation

$$E = T \cdot R,$$

where T is a skew-symmetric matrix, and R a rotation. The matrix T gives the translation in 3-space from the one camera to the other, and R their relative angular orientation. There is an ambiguity given by the alternative factorisation

$$E = (-T) \cdot (-R),$$

but this will not be of concern here.

Since E is a projective matrix, N = 8 sufficiently general point correspondences uniquely determine E. In fact, there exists a reconstruction algorithm which works in this way [16]. However, that method ignores the fact that an essential matrix is necessarily of rank 2. So, a 7-point algorithm came up replacing one of the equations in (3) by the cubic equation

$$\det(E) = 0$$

[9, 11]. A 6-point algorithm is described in [19]. It is known that the minimal number of point correspondences necessary for solving the relative pose problem is five. We now briefly review the essentials of the 5-point algorithm by [18].

The first step is to solve the linear system (3) with N = 5. The general solution is of the form

(4) 
$$E = x_1 E_1 + x_2 E_2 + x_3 E_3 + x_4 E_4,$$

where  $\{E_1, \ldots, E_4\}$  is a basis for the solution space of (3). Here, it is assumed that the 5 corresponding points are chosen in such a way that the rank of the system is five.

In the second step, the matrix (4) is plugged into the non-linear conditions for the essential matrix. These are given as

(5) 
$$2 \cdot EE^T E - \operatorname{Trace}(EE^T) \cdot E = 0$$

(6) 
$$\det(E) = 0$$

This yields 10 homogeneous cubic equations in the four unknowns x, y, z, w. The original method by Nistér is to set w = 1, eliminate variables and obtain a univariate polynomial f(z) of degree 10. The zeros of f(z) then lead to maximally ten candidate essential matrices (after discarding the non-real solutions).

Using the 2-adic image encoding from the previous subsection, we obtain for the 5-point relative pose problem the same equations (3), (5) and (6). The difference is now that the coefficients are 2-adic expansions of natural integers, and that the wanted solution is a set of 2-adic essential matrices. In the following subsection, we will explain how this can be obtained effectively by Hensel's lifting method.

4.3. Lifting the equations to  $\mathbb{Z}_2$ . The structure of the equations from the previous subsection depends on the particular sample of five corresponding pairs of points which in the following will simply be referred to as the *sample*. The algorithm later on will terminate either with a set of solutions or with a resampling routine, meaning that another set of five corresponding point pairs has to be chosen.

Let  $\mathscr{F} \subseteq k[x_1, \ldots, x_n]$  be a set of polynomials with coefficients in a field k. Then the zero set of  $\mathscr{F}$  will be denoted as  $V(\mathscr{F})$ :

$$V(\mathscr{F}) := \{ (\xi_1, \dots, \xi_n) \in k^n \mid f(\xi_1, \dots, \xi_n) = 0 \text{ for all } f \in \mathscr{F} \}.$$

This is also called a *variety defined over* k. Let R be a unitary commutative ring contained in the algebraic closure  $k^{\text{alg}}$  of k, and  $V \subseteq k^n$  a variety defined over k. Then the *R*-rational points of V are those points of  $V_{k^{\text{alg}}}$  lying in  $R^n$ , where  $V_{k^{\text{alg}}} \subseteq (k^{\text{alg}})^n$  is  $V(\mathscr{F})$  seen as a variety defined over  $k^{\text{alg}}$ . In particular, we will speak of k-rational points of a variety defined over k. The set of *R*-rational points of V will be denoted by V(R).

4.3.1. The linear equations. Assume that N = 5. The simplest case for applying Hensel's lifting method is that, after dividing off each equation the least common divisor of the coefficients, the linear equations (3) are linearly independent modulo 2. Then, by the multivariate linear Hensel lemma, e.g. [2, Thm. 2], there is a unique lift of a basis of the solution space of (3) modulo 2 to a  $\mathbb{Z}_2$ -basis of the solution space of (3). The constructive nature of the proof yields a lifting algorithm.

4.3.2. The cubic equations. The system (5) is modulo 2 of the form

(7) 
$$\operatorname{Trace}(EE^T) \cdot E \equiv 0 \mod 2$$

By construction, the entries of E are zero or homogeneous linear polynomials in  $x_1, x_2, x_3, x_4$ . As in 4.3.1, we assume that E is not the zero matrix modulo 2. In any case, the diagonal elements are zero or squares of linear forms. Hence, the polynomial

$$Q(x_1, x_2, x_3, x_4) = \operatorname{Trace}(EE^T) \mod 2$$

is a sum of squares:

$$Q = \sum_{i=1}^{4} \alpha_1 x_i^2 = L^2,$$

where  $\alpha_1, \ldots, \alpha_4 \in \mathbb{F}_2$  and

$$L = \sum_{i=1}^{4} \alpha_i x_i$$

is linear. From this it follows that (7) is of the form

(8) 
$$L^2 \cdot L_i = 0, \quad i = 1, \dots, 9,$$

with  $L_i$  either zero or linear modulo 2. If we assume that the  $\mathbb{F}_2$ -basis  $E_1, \ldots, E_4$  of (3) modulo 2 is read off a staircase normal form for the reduced linear system of equations, we observe that the matrix  $E \mod 2$  contains four entries consisting precisely of the variables  $x_1, x_2, x_3, x_4$ . Hence, the four equations  $L^2 \cdot x_i = 0$  are among (8), and it follows that the solution of that system is given by

$$(9) L = 0$$

over the finite field  $\mathbb{F}_2$ . However, the variety V(L) defined by (9) is a hyperplane in the projective space  $\mathbb{P}^3_{\mathbb{F}_2}$ , whereas (5) defines a curve in  $\mathbb{P}^3_{\mathbb{Q}_2}$  for a generic sample of five corresponding point pairs.

In any case, by the multivariate Hensel lemma [2, Thm. 1] the points of the set

$$V = \{x \in V(L) \cap V(\det(E) \mod 2) \mid \nabla \det(E)(x) \not\equiv 0 \mod 2\}$$

are uniquely liftable to  $\mathbb{Z}_2$ -rational points in  $\mathbb{P}^3$ , as long as det(E) is modulo 2 not the zero polynomial.

In order to obtain more liftable points, we take a closer look at the variety given by (5). Write to this aim

(10) 
$$E = (E_{ij}) = \begin{pmatrix} e_1 \\ e_2 \\ e_3 \end{pmatrix},$$

where  $e_i$  are the rows of E. Then (5) translates to

(11) 
$$(2EE^T - \operatorname{Trace}(EE^T) \cdot 1) \cdot E = 0.$$

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where 1 is the unity  $3 \times 3$ -matrix. By multiplying from the right with some invertible matrix, we may replace the rightmost matrix in (11) by a triangular matrix

$$T = \begin{pmatrix} T_{11} & T_{12} & T_{13} \\ 0 & T_{22} & T_{23} \\ 0 & 0 & T_{33} \end{pmatrix}.$$

Using (10), this is equivalent to the system

$$T_{11} \begin{pmatrix} e_1^2 - e_2^2 - e_3^2 \\ 2e_1e_2 \\ 2e_1e_3 \end{pmatrix} = 0$$

$$T_{12} \begin{pmatrix} e_1^2 - e_2^2 - e_3^2 \\ 2e_1e_2 \\ 2e_1e_3 \end{pmatrix} + T_{22} \begin{pmatrix} 2e_1e_2 \\ e_2^2 - e_1^2 - e_3^2 \\ 2e_2e_3 \end{pmatrix} = 0$$

$$T_{13} \begin{pmatrix} e_1^2 - e_2^2 - e_3^2 \\ 2e_1e_2 \\ 2e_1e_3 \end{pmatrix} + T_{23} \begin{pmatrix} 2e_1e_2 \\ e_2^2 - e_1^2 - e_3^2 \\ 2e_2e_3 \end{pmatrix} + T_{33} \begin{pmatrix} 2e_1e_3 \\ 2e_2e_3 \\ e_3^2 - e_1^2 - e_2^2 \end{pmatrix} = 0$$

where we use the sloppy notation  $e^2 = ee^T$  and  $e_i e_j = e_i e_j^T$ . This can be simplified to

(12) 
$$T_{11} \begin{pmatrix} e_1^2 - e_2^2 - e_3^2 \\ 2e_1 e_2 \\ 2e_1 e_3 \end{pmatrix} = 0$$

(13) 
$$T_{11}T_{22}\begin{pmatrix} 2e_1e_2\\ e_2^2 - e_1^2 - e_3^2\\ 2e_2e_3 \end{pmatrix} = 0$$

(14) 
$$T_{11}T_{22}T_{33}\begin{pmatrix} 2e_1e_3\\ 2e_2e_3\\ e_3^2 - e_1^2 - e_2^2 \end{pmatrix} = 0$$

Assume now that in (12)-(14) the least common divisor of the coefficients for each row in each equation (or just the highest common power of 2) has been divided off. The equations can now successively be simplified in a straightforward manner. This leads to a union of at most 14 varieties  $W_i$ , each of which is defined by quadratic and linear equations. We call this the sampled five-point trace variety W. Of interest is the case that  $V := W \cap V(\det(E))$  has dimension one.

Notice that

$$\det(E) = 0 \Leftrightarrow \det(A) = 0,$$

where  $A = (A_{ij})$ . We set

$$V_i := W_i \cap V(\det(A)) = V(\mathscr{F}_i) \subseteq \mathbb{A}^4,$$

where  $\mathscr{F}_i$  is the set of linear and quadratic polynomials from above, together with the cubic polynomial det(A). The sampled five points lead to finitely many candidate essential matrix if and only if the image  $\bar{V}_i$  under the canonical map

$$\mathbb{A}^4 \setminus \{0\} \to \mathbb{P}$$

is zero-dimensional for all *i* such that  $V_i \setminus \{0\} \neq \emptyset$ . The dimension can be checked on each affine chart  $x_i \neq 0$ . For this, we denote by

$$\mathscr{F}_i^j := \left\{ f|_{x_j=1} \mid f \in \mathscr{F}_i \right\}$$

and

$$V_i^j := V(\mathscr{F}_i^j) \subseteq \mathbb{A}^3$$

the affine piece of  $V_i$  given by  $x_j = 1$ .

If  $\mathscr{F}$  is a set of polynomials with coefficients in  $\mathbb{Z}_2$ , then  $\mathscr{F} \mod 2$  means the set consisting of the polynomials from  $\mathscr{F}$  with coefficients modulo 2. By  $V(\mathscr{F})$  mod 2 we mean the zero set of  $\mathscr{F} \mod 2$  defined over  $\mathbb{F}_2$ . We declare

$$\dim \emptyset := -1,$$

and arrive at the following lifting algorithm:

Algorithm 4.3. Input. The reduced matrix A with coefficients in  $\mathbb{Z}_2[x_1, x_2, x_3, x_4]$ . Step 1. Compute for all *i* the set  $\mathscr{F}_i$  as above, and  $\mathscr{F}_i \mod 2$ . If for some *i* the latter contains a non-zero polynomial, then continue. Otherwise, resample.

Step 2. Compute for all *i* the dimension  $d_i := \dim(\overline{V}_i \mod 2)$  on each affine piece  $V_i^j \mod 2$ ,  $j = 1, \ldots, 4$ . If all  $d_i \leq 0$ , then continue, otherwise resample.

Step 3. Compute all  $\mathbb{F}_2$ -rational points of  $V_i \mod 2$ , and for all such  $\omega \in V_i \mod 2$ the quantity  $\nabla f(\omega)$ , where  $f \in \mathscr{F}_i \mod 2$ . If some value is

$$\nabla f(\omega) \not\equiv 0 \mod 2,$$

then lift for that f and collect all lifts in the set  $\tilde{V}_i$ . If  $\tilde{V}_i \neq \emptyset$ , then continue. Otherwise, resample.

Step 5. Test for all i and all  $v \in \tilde{V}_i$  whether all quantities f(v) with  $f \in \mathscr{F}_i$  are zero. Collect all positively tested v in  $S \subseteq \mathbb{A}^4$ .

Output. The lifted finite set

 $\bar{S} \subseteq \mathbb{P}^3(\mathbb{Z}_2)$ 

of  $\mathbb{Z}_2$ -rational solutions.

Observe that there is a dimension computation in Step 2, and a solution set computation in Step 3, both for equations modulo 2. These can be effected with Gröbner basis methods, as described e.g. in [12, Cor. 3.7.26].

#### 5. p-adic random sample consensus via classification

In this section, we incorporate Algorithm 4.3 into a sampling scheme in which random samples of five point-pairs are taken, and the output of the lifting algorithm is collected, and then classified. The idea is that in the end, a pronounced cluster will appear in the classification which contains among its central elements the "true" essential matrix. Our approach differs from the original RANSAC [7] in that we fix the number of samplings instead of the cardinality of the consensus set, and that we perform a hierarchical classification of the solutions from each sample. The consensus set corresponds here to one of the clusters in the classification.

Let K be a p-adic field.

**Definition 5.1.** Let  $C \subseteq K$  be a cluster, and let  $A \subseteq C$  be the subset of all central elements with respect to E. Then the rooted tree  $\mathfrak{S}(C) := T^{\dagger}(A \cup \{\infty)\}$  is called the central spine of C.

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Since all central elements branch off the tips of  $\mathfrak{S}(C)$ , the central spine of a cluster C says something about the distribution of the data within C.

We define the *density* of a verticial cluster C as

$$\delta(C) := \begin{cases} \frac{|C| - 1}{\mu(C)}, & |C| > 1\\ 0, & \text{otherwise} \end{cases}$$

where

$$\mu(C) = \int\limits_{K} 1_{\mathbb{D}_C} \, dx,$$

with  $\mathbb{D}_C \subseteq K$  being the smallest disk containing C and

$$1_{\mathbb{D}_C} \colon K \to \mathbb{R}, \quad x \mapsto \begin{cases} 1, & x \in C \\ 0, & \text{otherwise} \end{cases}$$

the characteristic function.

The following algorithm is a p-adic analogon of a variation of the Ran(dom) Sa(mple) C(onsensus) algorithm [7] applied to the problem of estimating the essential matrix from two images. In this variation, the consensus is established by a hierarchical classification of the collected solutions for the equations given by the sampled five-tuples of corresponding image points. In order to establish the "winning" cluster, we consider each sample as casting upto ten votes. Then we establish a ranking of clusters according to the following criteria:

- (1) majority of votes
- (2) highest density
- (3) highest precision

These criteria are to be taken in that order, i.e. the clusters are ranked according to criterion (1). Ties are first broken using criterion (2), and then with criterion (3). This means a ranking of clusters

- (1) according to their size
- (2) according to  $\delta(C)$
- (3) according to  $\mu(C_c)$ ,

where the  $C_c$  is given by the following definition:

**Definition 5.2.** Let  $C \subseteq K$ . Then the central cluster in C is the smallest verticial subcluster of C containing the centres of C.

**Remark 5.3.** In the case that the differences in size, density or precision are small among the high ranked clusters, it makes sense to allow almost equally ranked clusters to be considered as ties and to use the next criterion to break them.

Fix an isometric isomorphism  $(K, |\cdot|_K) \cong (\mathbb{Q}_2^{3 \times 3}, \|\cdot\|_{\max})$ , where K is an unramified extension field of  $\mathbb{Q}_2$ , and  $\|\cdot\|_{\max}$  is the maximum of the 2-adic norms of all matrix entries.

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**Algorithm 5.4** (RanSaC<sub>p</sub>). Input. Numbers  $k, m, n, N \in \mathbb{N}$  and a finite set  $X \subseteq I \times I'$  of pairs of corresponding points in two images I and I' represented as 2-adic vectors.

Step 1. Sample five random elements of X. If the equations (3) modulo 2 have rank 5, then solve these by lifting a basis modulo 2 to a basis modulo  $2^n$ . If  $n \gg 0$ , then the lift yields exact solutions in  $\mathbb{Z}_2$ . If modulo 2 the rank is smaller than five, then resample.

Step 2. Perform Algorithm 4.3 by lifting to solutions modulo  $\mathbb{Z}/2^m\mathbb{Z}$ , where m is the desired precision. Obtain a set of approximate candidate essential matrices with entries in  $\mathbb{Z}/2^m\mathbb{Z}$ . If that set is non-empty, then continue. Otherwise, resample.

Step 3. Repeat Steps 1 and 2 successively N times, and obtain an accumulated set  $\mathcal{E}$  of approximate candidate essential matrices from each repetition.

Step 4. Use the  $LBG_p$  algorithm over K for obtaining  $\leq \ell$  clusterings of  $\mathcal{E}$  with  $\ell = 2, \ldots, k$ . Determine the ideal clustering(s), in the sense of Definition 3.5, within  $\mathfrak{X}_k$ . In these, determine the winning clusters, their centres and central spines.

Output. A set of approximate central essential matrices.

**Remark 5.5.** The desired result of an instance of  $\operatorname{RanSaC_p}$  applied to genuine image data would be a clustering in which there is one pronounced cluster C having a central spine which is a path segment of length n >> 0. In this case, the central elements of C would yield one single candidate essential matrix  $E \in (\mathbb{Z}/2^n\mathbb{Z})^{3\times 3}$ approximating the "true" 2-adic essential matrix for the particular stereo image problem, while the other clusters could be considered as "outliers". In general, noise in the image will lead to less pronounced clusters and central spines with branching. The former can lead to wrong estimates for E, and the latter means that the approximated essential matrix can be less precise than in an ideal setting.

## 6. CONCLUSION

The *p*-adic classification algorithm  $\text{LBG}_p$  is incorporated into a random sample consensus algorithm via classification  $(\text{RanSaC}_p)$  in order to efficiently solve the five-point relative pose problem in stereo vision. The equations occurring in the relative pose problem are derived from a 2-adic encoding of image coordinates, decomposed and then solved with Hensel's lifting method. The cluster number is determined with a *p*-adic version of an intra-inter-validity measure originally developped for *k*-means. The proposed solution for the essential matrix lies in the centre of the most significant cluster.

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