

Force Field Simulation Based Laser Scan Alignment

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1. Introduction

Alignment of sensor data, typically acquired from cameras, laser range scanners, or sonar sensors, is the basis for all robot mapping tasks. Recent advances in the development of laser range devices make research on laser range alignment a focus of robot mapping research. In contrast to cameras, laser range scanners offer relatively precise depth information, yet the feature density is relatively sparse. Since alignment algorithms are based on feature correspondence, a lack of features naturally causes problems. One way to approach that problem is to cover the area with a high number of scans, such that subsequent scans have a low relative displacement only. This guarantees sufficient scan overlap and a reliable detection of feature correspondences. Though this approach is feasible for many mapping applications, it can not be assumed for an important field of robotics, namely Urban Search and Rescue Robotics ("rescue robots"), in there especially the setting of multi robot mapping. In multi robot mapping, a number of robots scan the environment independently, without reliable knowledge of their relative position. Additional sensors, like GPS, can not be assumed due to the nature of the environment. Non autonomous rescue robots were e.g. deployed after the 9/11 attack to assist in the search for victims in the collapsed towers. In such an environment, GPS is not available because of the massive concrete walls surrounding the robots. The task of multi robot mapping in rescue environments imposes especially challenging constraints:

- no precise or reliable odometry can be assumed, which means especially that the robots' relative poses are unknown
- due to the nature of catastrophe scenarios no distinct landmarks are given
- the overlap between pairs of the robots' scans is minimal

Figure 1 shows 12 out of 60 single scans from multiple robots, taken in a disaster test area at NIST, Gaithers-burg, MD. Even for humans it is hard to detect overlapping features.

Our approach to alignment of such a data set is to first give a rough estimate of the robots' poses, called the pre-alignment, and then to improve the achieved map. This article deals with the second step, the improvement, see figure 2.

This article introduces of a new process, called 'Force Field Simulation' (FFS), which is tailored to align maps under the aforementioned constraints. It is motivated by simulation of dynamics of rigid bodies in gravitational fields, but replaces laws of physics with constraints derived from human perception. It is an approach of the family of gradient

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descent algorithms, applied to find an optimal transformation of local maps (in particular, laser range scans) to build a global map based on feature correspondences between the local maps. Figure 3 shows the basic principle: forces (red arrows) are computed between 4 single scans (the 4 corners). The scans are iteratively transformed by translation and rotation until a stable configuration is gained. The single scans are not merged, but kept separated. As they are moved according to the laws of the motion of rigid bodies in a force field, single scans are not deformed. FFS has the following properties:

1. Low level correspondences (data point correspondences) are not made by a hard decision (an integral of forces between pairs of points defines the force field in place of hard 'nearest neighbor' correspondences)
2. FFS is a gradient approach, it does not commit to an optimal solution in each iteration step
3. The iteration step towards an optimal solution is steered by a 'cooling process', that allows to jump the system out of local minima
4. FFS transforms all scans simultaneously
5. FFS easily incorporates structural similarity modeling human perception to emphasize/strengthen the correspondences

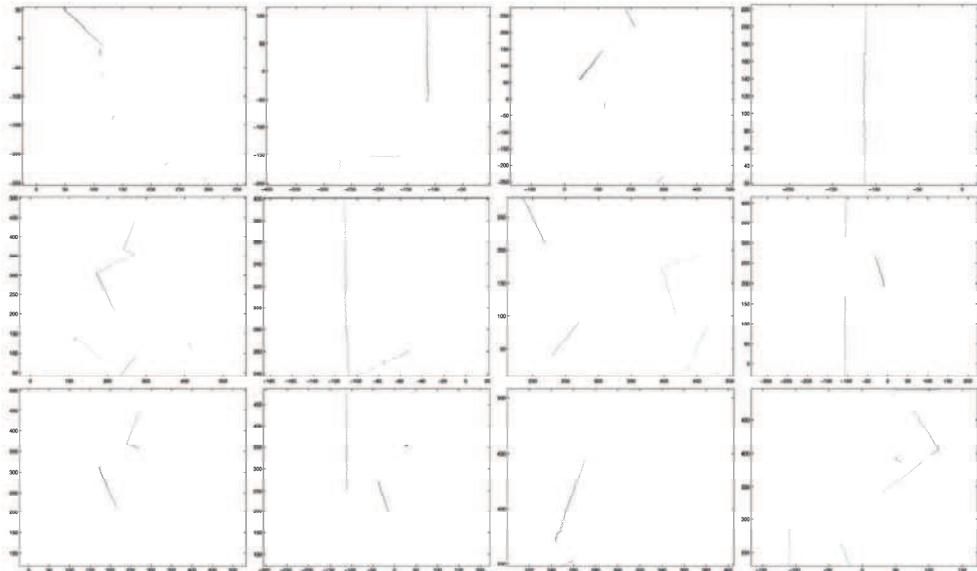


Figure 1. 12 out of 60 single scans from the NIST disaster test area. Even for humans it is hard to detect overlapping features

2. Related work

The problem of aligning n scans has been treated as estimating sets of poses (Lu and Milios, 1997). Since sets of poses and the associated structures (maps) are conditionally independent, this estimation is Simultaneous Localization and Mapping. The conditional independence is e.g. the key for Rao-Blackwellization (factoring the posterior of maps) of particle filters for SLAM (Montemerlo et al., 2002).

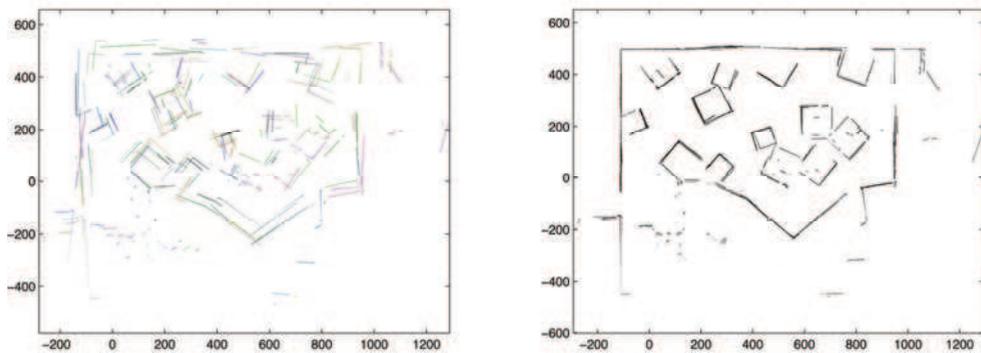


Figure 2. Left: Pre-alignment of the 60 scans of the NIST disaster test area. Right: improved alignment. Force Field Simulation is used to achieve the improved alignment, given an estimated pre-alignment

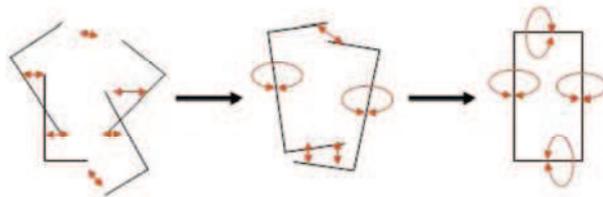


Figure 3. Basic principle of FFS. Forces are computed between 4 single scans. Red arrows illustrate the principle of forces. The scans are iteratively (here: 2 iterations) transformed by translation and rotation until a stable configuration is achieved

There have been several algorithms to estimate the sets (Olson et al., 2006; Frese, 2006; Frese et al., 2005; Thrun et al., 2002; Minguez et al., 2006; Konolige, 2003). The underlying framework for all such techniques is to optimize a constraint-graph, in which nodes are features, poses and edges are constraints built using various observations and measurements like odometry scan-matching of range scans. These techniques differ in

- how they represent graphs - e.g. (Frese, 2006) uses a sophisticated data structure called Tree-map, (Thrun et al., 2002) represents using sparse extended information filters (SEIF).
- how they build constraints - e.g. (Lu and Milios, 1997) uses linearized constraints obtained from scan-matching and odometry, (Olson et al., 2006) works with non-linear constraints.
- how they optimize the graphs - e.g. (Olson et al., 2006) uses stochastic gradient descent for approximate optima, borrowing the ideas from learning theory, (Lu and Milios, 1997) solves for exact optima using brute-force, (Frese et al., 2005) use Gauss-Seidel relaxation again for approximate optima.

All these approaches have performed well in many practical cases but they have one drawback that is they are sensitive to behavior of error models of sensors because of several assumptions and approximations which might not hold with sparse sensing.

(Lu and Milios, 1997) linearizes constraints by linearizing pose-relations, solving a linear equation of the form $AX = B$ to estimate X , the set of poses. This requires that A is invertible, so they conjecture that A is invertible if the constraint-graph is fully connected and the errors of the observations behave in a gaussian/normal way. (Borrmann et al., 2007) extends the same technique for 3D scans.

(Olson et al., 2006) presents an approximate optimization of non-linear constraints and demonstrate that their approach of approximating the optimization process in non-linear state space yields superior results compared to finding exact optima by approximating a non-linear state space (SLAM) to a linear state space.

Another strategy of attacking the problem is to treat the problem of SLAM from a perspective of aligning n scans simultaneously. The algorithms exploiting this perspective build from image registration techniques, the most famous being Iterative Closest Point (TCP) (Besl and McKay, 1992; Chen and Medioni, 1992) and its numerous variants to improve speed and converge basins (Rusinkiewicz and Levoy, 2001) and (Lu and Milios, 1994; Birk and Carpin, 2006). Basically all these techniques do search in transformation space trying to find the set of pair-wise transformations of scans by optimizing some function defined on transformation space. The techniques vary in defining the optimization functions that range from being error metrics like "sum of least square distances" to quality metrics like "image distance" as in (Birk, 1996). Their optimization process itself can be gradient descent or hill climbing or using genetic programming strategy as in (Robertson and Fisher, 2002). All of these techniques have one major limitation, which is they search in *pair-wise* transformation space. Though in some variants of ICP the error from all pair-wise transformations is spread across all transformations to simultaneously align all scans, the procedure can be highly sensitive to outliers (Rusinkiewicz et al., 2005).

FFS also adapted the perspective of aligning n scans, it treats the alignment problem as an optimization problem. Rather than using a least squares solution to compute intermediate motions, FFS uses an iterative gradient technique to solve for (local) optima. Here FFS is similar to the approach proposed by (Eggert et al., 1998), which simulates a dynamic spring system to register multiple range scans simultaneously. They describe the advantages of such a gradient descent system as follows: *'The reason [not to use a least square solution] is that the effects of any significantly incorrect correspondences are compounded when the best alignment is computed (...) With a dynamic system it is possible to move in the direction of an intermediate solution without being totally committed to it'*. (Eggert et al., 1998) differ in the choice of the registration function, which in contrast to FFS is based on one to one correspondences between points, as well as in the optimization technique.

FFS uses a gradient method with decreasing step width Δ_i . The registration function (target function) of FFS is based on Gaussian fields, similar to (Boughorbel et al., 2004). In contrast to (Boughorbel et al., 2004), FFS uses a variable, decreasing σ for each iteration step t . Additionally, (Boughorbel et al., 2004) solve the optimum of the registration using a quasi-Newton method, hence they do not steer the system with a step width parameter.

Since we keep the single scans separated, our search space is high dimensional, in the 2D case it is $3n$ -dimensional (3D: $6n$ -dimensional), with n being the number of scans. For example, our experiment described in section 4.2 uses 60 scans, our search space is therefore 180-dimensional. Birk and Carpin use a random walk technique to reach the optimal solution. Since random walk techniques tend to become critical in high dimensions, we do not utilize this technique in our approach but decide in favor of a guided (gradient) walk.

This search in high dimensional space at first sight seems very complicated, demanding computation of high dimensional gradient but fortunately using potential field simulation for various computer vision tasks like contour detection, segmentation, registration has been empirically successful (Yang et al., 2006; Jalba et al., 2004; Xu and Prince, 1998), (Ayyagari et al., 2005), (N.Paragios et al., 2003), (Veltkamp and Hagedoorn, 1999). Since mapping is closely related to registration, the approaches whose motivations are closely related to our approach are (Biber and Strasser, 2006; Ayyagari et al., 2005; Eggert et al., 1998). In (Eggert et al., 1998) they align range scans by moving them simultaneously. The movements are not just based on the minimizing error of transformation computed using correspondences but on the simulated fields generated by imaginary springs attached to the corresponding points. Our technique differs from (Eggert et al., 1998) in that the force field is generated not just by closest point correspondences but using perceptual principles and gaussian fields similar to (Boughorbel et al., 2004). (Biber and Strasser, 2006) also performs search in 3n dimensional space. For each configuration they compute energy as the sum of the Normal Distribution Transforms (NDT) (Biber and Strasser, 2003) of all the scans in the configuration and update the configuration using Newton's optimization algorithm that involves the first and second derivatives of the energy. Their approach is very closely related to ours but does not use perceptual features and rigid body dynamics and hence in principle can be more sensitive to outliers.

3. Force field based mapping

The following motivates and describes the FFS algorithm. A pseudo code representation can be found at the end of this section.

3.1 Basic Principle

To draw the analogy to Newtonian Physics, each scan s_i can be seen as a rigid body of masses: the scan points represent the masses, rigidly connected by massless rods. A global map g defines the transformation of all scans, it therewith defines the distribution of all masses (the union of all scan points). In the framework of Newtonian Physics the gravitational forces between these masses forms a gradient field. The FFS algorithm is motivated by simulation of the movement of bodies in a gradient field. In contrast to pure physics it replaces physical principles of masses and forces by principles that correspond to human visual perception, i.e. gravitation is replaced by 'strength of correspondence'. Also, to achieve convergence to a stable state of minimal total energy, the kinetic energy is not taken into account, i.e. the velocity of each rigid body after each iteration step is set to 0. Also FFS uses a 'cooling' strategy in its step width parameter that initially adds energy to the system to allow for escape from local minima, see section 3.2.

Let $S = s_1, \dots, s_n$ be a set of n scans gained from laser range scan devices. A scan $s_i = (p^i_1, \dots, p^i_j)$ consists of j data points. Data points are the coordinates of reflection points of the laser range scanner in a local coordinate frame defined by a single robot pose. We also assign a scalar value, a *mass* m^i_j , to each data point, which can be interpreted as the perceptual importance. For the purpose of multi robot mapping we assume that each scan is possibly gained from a different robot, while the robots' relative poses are unknown or poorly estimated. The task of the algorithm is an optimization over the set of the robots' poses; hence the goal is to find transformations for all n scans $s_{i=1..n}$ to registrate the scans, such that

similar features in different scans match 'perceptually consistently' when they are superimposed on top of each other.

Observe that the order of local maps is irrelevant in our framework since we transform the scans simultaneously, which is an important property of the algorithm to be applicable for multi robot mapping. For single robot mapping, FFS is canonically extendable to online FFS, see section 3.5.

The transformations performed are rotation θ_i and translation x_i, y_i of each scan s_i . Superimposing the transformed maps builds a *global map* g as shown in Fig. 4.

During the global map building process single maps are not merged but kept separated. Therefore a global map is defined by the vector of the transformation parameters of all scans: a *global map* $g = (t_1, \dots, t_n)$ is a $3n$ -dimensional vector of transformation parameters $t_i = (x_i, y_i, \theta_i)$ of all n scans $s_{i=1..n}$. The space of all global maps is denoted by \mathcal{G} . To register the scans, we define a *fitness measure* P_g to evaluate the 'perceptual consistency' of a global map g . Finding the global map g_k that minimizes P_g is clearly an optimization problem. FFS solves this optimization problem with a gradient technique that iteratively transforms all scans simultaneously until a stable configuration (local minimum) is reached. The following section will motivate and define the fitness measure P_g as well as the implementation of the gradient approach.

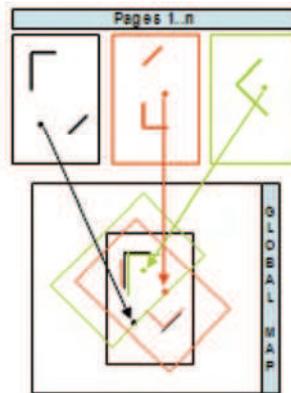


Figure 4. Single scans are transformed to build a global map

3.1.1 Correspondence function

As in (Boughorbel et al., 2004), the basic idea of our registration method is to use a Gaussian field to define a strength of correspondence between data points, i.e. a measure for both spatial proximity and visual similarity of two points belonging to different scans.

A *correspondence* between data point p_1 and a data point p_2 is defined as a vector

$$V(p_1, p_2) = C(p_1, p_2) \frac{p_2 - p_1}{\|p_2 - p_1\|} \quad (1)$$

Its magnitude $\|V(p_1, p_2)\| = C(p_1, p_2)$ describes the strength of correspondence, defined as:

$$C(p_1, p_2) = \frac{1}{\sigma_t \sqrt{2\pi}} e^{\left(-\frac{\|p_2 - p_1\|^2}{2\sigma_t^2}\right)} m_1 m_2 \cos(\angle(p_1, p_2)) \quad (2)$$

with m_i being the mass assigned to p_i , and the angle $\angle(p_1, p_2)$ being the angle between the directions of points p_1, p_2 , which will be defined in section 3.3. Intuitively, the direction of a point is the direction of an underlying model of a linear structure (a line segment).

A major difference to the pure physics simulation is that the mass values assigned to the data points are not assumed to be constant. The mass m_i for a point p_i is used to compute the force as in eq. 3, yet it can be reassigned a different value for the computation of movement of the scan (we are not modeling physics but perception, hence freedom from Newton's law is given). Steering the mass enables the algorithm to react better to perceptual properties: there is no perceptual reason for an 'important point', e.g. a corner point, assigned a high mass for force computation, to be less mobile than other points during movement computation (caused by its high mass). This observation suggests using different masses during the computation of forces than during the computation of the movement. In the current system, the mass assignment is used to regulate influences of different scan densities, i.e. high density points (due to a high number of scans in a single area) are assigned lower mass values, while low density points (above a certain threshold) are slightly overemphasized using higher mass values.

Expression 2 can be interpreted as a force field whose sources are located in the data points. It has the following properties:

1. The strength of correspondence decays with Euclidean distance, the influence of distance is controlled by the parameter σ_t
2. The strength of correspondence is weighted by the mass of each data point and depends on the angle between point directions, i.e. it is 0 for orthogonal directions, 1 for parallel directions.

We propose this model for the following reasons:

1. Distance likelihood and parallelism follow the basic principles of Gestalt Psychology (Wertheimer, 1958), modeling low level cognition.
2. The scale parameter a_t gives additional freedom to adjust the process (see section 3.2), it enables the correspondence process to work on different visual scales.
3. Assigning a mass to a data point can be seen as assigning a visual importance to it. Data points in higher regions of interest can be stronger emphasized.

In terms of the physical framework, the correspondence $V(p_1, p_2)$ (eq. 1) describes a force on p_1 towards p_2 with strength $C(p_1, p_2)$. Embedding the scans s_i into \mathbb{R}^2 using the transformations defined by a global map g , we can define a vector field $F: \mathbb{R}^2 \supset \mathcal{P} \rightarrow \mathbb{R}^2$, the *Force Field* on the set of all points $\mathcal{P} = \{p | p \in \bigcup_{i=1..n} s_i\}$ by summing the correspondences.

$$F(p_i) = \sum_{p_j \in \mathcal{P} \setminus p_i} V(p_i, p_j) \quad (3)$$

By definition of the strength of correspondence F is radial and hence a gradient field. With

$$A = m_1 m_2 \cos(\angle(p_1, p_2))$$

the overlying potential is defined by

$$P(p_i) = \frac{1}{2} \sum_{p_j \in \mathcal{P} \setminus p_i} \int_{-\infty}^r \frac{A}{\sigma_t \sqrt{2\pi}} e^{\left(-\frac{z^2}{2\sigma_t^2}\right)} dz \quad (4)$$

with $r = \sqrt{(X-x)^2 + (Y-y)^2}$, $p_i = (X, Y)$, $p_j = (x, y) \in \mathcal{P}$.

Note: $P(p_i)$ is the potential over F since:

$$\begin{aligned} F(p_i) &= -\nabla P \\ &= \begin{bmatrix} -\frac{\partial P}{\partial x} \\ -\frac{\partial P}{\partial y} \end{bmatrix} = \begin{bmatrix} \sum \frac{A}{\sigma_t \sqrt{2\pi}} e^{\frac{-r^2}{2\sigma_t^2}} \cdot \frac{X-x}{r} \\ \sum \frac{A}{\sigma_t \sqrt{2\pi}} e^{\frac{-r^2}{2\sigma_t^2}} \cdot \frac{Y-y}{r} \end{bmatrix} = \sum \frac{A}{\sigma_t \sqrt{2\pi}} e^{\frac{-r^2}{2\sigma_t^2}} \cdot \vec{u} = \sum V(p_i, p_j) \end{aligned}$$

where $\vec{u} = \frac{p1-p2}{\|p1-p2\|}$ and all sums are defined over $p_j \in \mathcal{P} \setminus p_i$.

Finally, we define the *fitness measure* or *potential* of a global map $g \in \mathcal{G}$ as the sum of potentials of all data points $p \in (P)$:

$$P(g) = \sum_{p_i \in \mathcal{P}} P(p_i) \quad (5)$$

In this framework, the potential $P(g)$ can be interpreted as the weighted average distance of all point pairs of different scans, the weight being the strength of correspondence. This potential can be seen as the quality of registration achieved by the transformations defined by g . To minimize the potential we apply an iterative gradient descent approach, the gradients in each data point given by $F(p_i)$, eq. 3. Computing the correspondences explicitly gives us these gradients, hence in the implementation of the algorithm there's no need to explicitly compute and derive the potential $P(g)$ (eq. 5) for the actual gradient descent.

In FFS, the computation of the transformation of each scan is determined assuming movement of rigid bodies in the given gradient field, i.e. all data points $p_i^j \subset \mathcal{P}$ of a single scan s_i share the same transformation, consisting of rotation and translation. However, eq. 3 does assume a non-rigid, independent movement of the data points, also the points' potential $P(p_i)$ (eq. 4) is defined over the space of all single (not rigidly connected) point configurations, which is a $2m$ dimensional space with $m = |\mathcal{P}|$. This means, the gradient $F(p_i)$ is defined under the assumption of unrestricted freedom of movement in \mathbb{R}^2 . To implement rigid body movement, we have to impose a restriction on the movement. The restriction is defined by the possible point configurations that are allowed by the transformations $g \in \mathcal{G}$. The laws of rigid body dynamics define these constraints: computing the gradient $F(p_i)$ in each data point as in equation 3 results in a $2m$ -dimensional gradient vector, the laws of rigid body dynamics map this vector to a $3n$ dimensional vector ∇g . ∇g describes the transformation of all n scans such that each data point moves in the direction of maximum descent of $P(g)$ in eq. 5, i.e. under the rigid body constraints. We therefore achieve a movement in the direction of the gradient gained in the $2m$ dimensional single point space projected onto the restricted $3n$ dimensional rigid movement subspace.

The basic laws of dynamics of rigid bodies in force fields accumulate the translation of all masses of a single scan into a single translation and a single rotation around a defined center. For each scan s_i the translational and rotational acceleration has to be determined. The translational acceleration $\alpha_T(s_i)$ of a scan s_i is defined by:

$$a_T(s_i) = \frac{\sum_{p \in s_i} F(p)}{\sum_{p \in s_i} m_p} \quad (6)$$

The rotational acceleration α_R is computed by torque and moment of inertia. Torque and inertia play the role of force and mass respectively, but take into account the distance to the rotational center c_R .

$$\begin{aligned} inertia &= \sum_{p \in s_i} m_i \|p_i - c_R\|^2 \\ torque &= \sum_{p \in s_i} \|p_i - c_R\| \times F(p) \end{aligned}$$

α_R is defined as

$$a_R = \frac{torque}{inertia} \quad (7)$$

The rotational center c_R is either defined as the robot's position, or by the center of mass. Experiments show, that in the first iteration steps it is useful to set the rotational center to the center of mass, while in later steps the robot's position is preferable. The first choice enables easier rotation, the second is modeling the actual scan setting more precisely. Hence, the closer the global map is to the solution, the more preferable is the robot's position as rotational center.

With α_T and α_R the transformation $t_k = (x_k, y_k, \theta_k)$ for scan s_k is defined by:

$$(x_k, y_k) = \frac{1}{2} a_T \Delta_t^2 \quad (8)$$

$$\theta_k = \frac{1}{2} a_R \Delta_t^2 \quad (9)$$

Δ_t being the step width of the gradient descent, as described in section 3.2.

With these constraints, the gradient in each iteration is computed by the following steps:

1. for each pair of points $p_i, p_j \in \mathcal{P}$ compute $V(p_i, p_j)$
2. for each point $p_i \in \mathcal{P}$ compute $F(p_i)$
3. for each scan $s_i \in S$ compute the transformation $t_k = (x_k, y_k, \theta_k)$ using the points $p^k \in s_k$.

This step results in a $3n$ dimensional gradient vector ∇g .

Computing all correspondences V in step 1 is an $O(n^2)$ process, section 3.4.1 will deal with the necessary reduction of computational complexity.

Figure 5 shows 2 iteration steps of FFS using 2 simple scans, consisting of 3 and 5 data points. In the left figure, the forces $F(p_i)$ in each data point p_i are plotted as green dotted lines. The 2 scans are transformed until they are superimposed, i.e. a stable configuration (local minimum of $P(g)$) is reached.

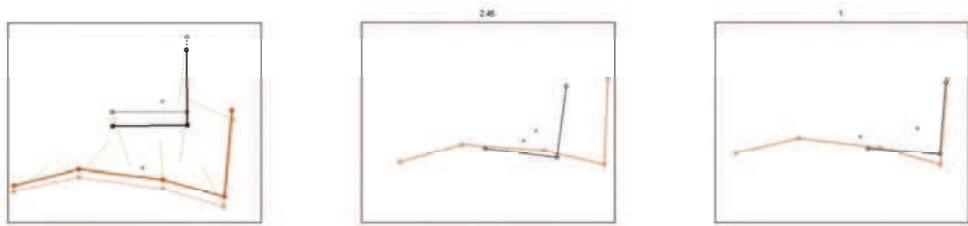


Figure 5. Left: Two scans (black/brown) superimposed. Dotted lines: scans at time t , green lines: forces on the data points at time t . Solid lines: scans after one iteration, time $t + \Delta t$. Both scans are translated/rotated according to the forces. Center: after iteration 5. Right: iteration 10

As in all gradient descent methods, the determination of the step width Δ_t is crucial. Also, gradient methods imply the danger of being trapped in local minima. We tackle both problems with the determination of step with Δ_t and σ_t as described in the following section.

3.2 Cooling down the motion: time stepping Δ_t and parameter σ_t

The determination of step width parameter Δ_t in any gradient descent approach is a well known problem. At chosen too small results in inapplicable slow convergence behavior and is not robust too noise, Δ_t chosen too big might miss the optimum. In FFS, the step width Δ_t is used as a steering parameter of the algorithm in connection with the parameter σ_t which determines the influence of distance in the correspondence function. We designed Δ_t as exponentially decreasing, σ_t linearly decreasing.

A large Δ_t allows the scans to be massively relocated (shuffled), they overshoot their correct position in the direction of the correspondence gradient. Naturally, a small Δ_t moves the scans less (the amount of replacement is directly proportional to Δ_t^2 , as defined by the laws of movement). We chose the strategy of decreasing Δ_t and σ_t experimentally, having analogies of the cooling behavior of algorithms like simulated annealing in mind. The imprecise, non optimal large Δ_t at the beginning allows the system to possibly escape from local minima. Observe that in contrast to a technique like simulated annealing we cool down a gradient guided process, not a random state change or a random walk technique that would not be applicable in our high dimensional search space. We therefore avoid the problems with a high computational load (high number of iteration steps) that tend to appear in simulated annealing due to unguided selection of the next state.

The parameter σ_t in equation 2 steers the influence of distance in the computation of point correspondences. A large σ_t enhances the relative influence of data correspondences with greater distances, and, since it equalizes this spatial proximity property, favors the influence of visual similarity. A small σ_t emphasizes local proximity, which is useful if the global map is already close to an optimum.

The effect of cooling is shown in figure 11 showing our experimental results on the 'apartment data set'. Observe that the potential function (the fitness measure) 12 is not monotonically decreasing in the first iteration steps. This shows an 'overshooting' of the system due to a large Δ_t , in the data registration this can indicate an escape from a local minimum.

It is important to mention that after each iteration the system resets the velocity of each scan to zero. This guarantees that the system converges to a stable state (assuming $\Delta_t \rightarrow 0$).

3.3 Point direction and optional resampling

The 'point direction' is used in the correspondence equation 2 to assign to points the direction of an underlying linear structure. It is derived by modeling the point set with line segments using the extended EM algorithm described in (Latecki and Lakaemper, 2006). Utilizing a segment split and merge approach, the extended EM algorithm automatically adjusts number and location of the line segments in a way such that linear structures are represented (by a single, possibly long line segment) as well as round structures (by multiple short segments). Hence, even scenarios not being rich in linear structures are robustly represented. The algorithm was already successfully applied to model indoor and outdoor rescue scenarios. A 3D version of this algorithm for approximation of scan points with planar patches is described in (Lakaemper and Latecki, 2006).

The data of each scan s_i is approximated by a set of line segments L_i . The direction of a point p in s_i is the angular direction (in the scan's coordinate frame) of the closest line segment in L_i . The approximation of the data set with line segments results in a very stable and intuitive estimation of point directions. Figure 6 shows the influence of point directions for the correspondences.

The closest line segment l_j to the point p is called the *supporting line segment* if its distance to p is below a certain threshold. Points without supporting line segments are removed from the data set. Due to the nature of the extended EM algorithm, these removed points are points in areas with low point density. Low point density results from objects which are hit less than others. This is the result of either erroneous scanning, non static objects, or low scanning density, which by itself results from either long distance to an object or simply the fact that a certain location was only scanned a few times. All of these topics include uncertainty about the existence of the object, hence we disregard such points. The behavior of the extended EM segment fitting guarantees that safe, distinct objects are not removed. The removal of uncertain data increases the stability of the FES algorithm.



Figure 6. Left: forces between two scans (red lines belong to first scan, black lines to second scan) computed without direction information. Right: forces computed using L and K as described in example 2 (using equal masses). Correspondences between non parallel structures are weakened

Having the segments, the data can optionally be resampled along the supported line segments with an equal sampling distance. Such a point set has a more homogeneous distribution of points, which tends to be advantageous: experiments showed that homogeneous distributions are helpful to avoid local minima if the configuration of scans is

still far from the optimal solution, since over represented areas (e.g. features with unusual high scanning density due to multiple scans in a single location) are equalized. Additionally, the optional resampling can significantly speed up the computation if the number of data points reduces drastically due to the chosen sampling resolution, see section 3.4.1. If the data is resampled, only the line segments (two endpoints) are stored, also resulting in a significant data compression (typically about 1:100).

3.4 Computational complexity

3.4.1 Time complexity

The definition of C (equation 2) on pairs of data points leads to an algorithm with $O(n^2)$ time complexity where n is the number of points. This is certainly prohibitive for real applications. Different techniques can be used to reduce the complexity by taking advantage of two main properties of equation 2:

1. For each data point only its local neighborhood must be examined, since the forces between points rapidly decrease with distance. Hence some techniques successfully built into ICP implementations (which suffers from the same complexity problem) can be used to reduce the complexity, e.g. KD-trees. In the current implementation, we take advantage of the line segment representation of the data; we use a bounding box intersection approach on axis aligned bounding boxes around the line segments: bounding boxes around all line segments are computed and extended by $2 \sigma_i$ in each direction. The force between two data points is computed only if the two corresponding lines' bounding boxes overlap. Hence we first reach a computational complexity based on the number $m \ll n$ of line segments, which is significantly lower than the number n of data points. Secondly, though bounding box intersection is an $O(m \log m)$ computation (note again: m =number of segments), update techniques as reported in (Cohen et al., 1995), reduce the expected complexity to $O(m)$. This linear complexity is reported under the constraint of 'relatively small' movements of objects, such that the $O(m \log m)$ sorting in the sweep and prune step reduces to $O(m)$ on a nearly presorted list. The constraint of small movements is met for most of the iteration steps in FFS. To give an idea of the order of magnitude of reduction that is achieved some numbers for the NIST data set (see section 4.2) should be mentioned:
 - 60 scans contain a total of 21420 points, represented by 332 line segments (on average: 65 points per segment)
 - average number of colliding pairs of segments per iteration 1500, hence we have $65 \times 65 \times 1500 (= 6,337,500)$ computations, compared to $21420^2 (= 460,000,000)$.
2. The data points have to be evaluated with a certain accuracy only. By approximating the evaluation of force field we can achieve computational reduction in the following 2 manners:
 - The current FFS implementation sub samples each segment equally with some sampling distance. For the NIST data set (sampling distance 10cm), we achieve in average 7 data points per segment, the force computation is therefore reduced to $7 \times 7 \times 1500 (= 73,500)$ computations.
 - Greengard and Strain introduced Fast Gauss Transform (FGT) (Greengard and Strain, 1991) which is in turn based on Fast Multipole Methods introduced for high speed simulation of particle dynamics in potential fields (Greengard and Rokhlin, 1987). The main advantage with FGT is that the force field can be computed in

linear time with a constant factor depending on the precision required in computation of the field. Details can be found in (Greengard and Strain, 1991). The main idea is to compute the force field using a divide and conquer strategy and exploiting Hermite and Taylor expansions. FGT has been first introduced in (Elgammal et al., 2003; Ayyagari et al., 2005).

3.4.2 Space complexity

Since we approximate the scans by segments, it is not necessary to keep the original data. For each scan, only the segments' endpoints have to be stored. Experiments with the extended EM algorithm (Latecki and Lakaemper, 2006) on 2D Laser data sets show an average compression rate of 1:100 (200 data points per segment).

3.5 Online FFS

The described algorithm easily can be extended to online SLAM, i.e. scans are recorded and processed subsequently, as they arrive from the laser device. The extension is canonical: each additional scan is pre aligned, then FFS runs on the previously aligned data set plus the new scan. The current FFS system targets the application of multi robot mapping, hence the sequential processing is not implemented yet.

Algorithm 1 Force field based mapping

- 1: Compute supporting line segments (section 3.3)
 - 2: Resample data set (section 3.3)
 - 3: $S \leftarrow$ initial state of transformations
 - 4: initialize step width Δ_t and σ_t (section 3.2)
 - 5: **repeat**
 - 6: Assign masses to data points
 - 7: Compute forces using σ_t (eq. 3)
 - 8: Assign constant mass values to data points
 - 9: Compute rotational and translational acceleration (eq.7 and eq. 6)
 - 10: Compute transformations ∇g
 - 11: $g \leftarrow g + \nabla g$
 - 12: Set velocity of all points to zero
 - 13: update σ_t and Δ_t (section 3.2)
 - 14: **until** average of relocations > Threshold
-

3.6 Pre Alignment

The pre alignment does not make use of odometric sensor data, but is based on shape similarity. It finds distinct shape features in single scans and tries to find an optimal overlap based on the shape similarity of these features. For further details see (Adluru et al., 2006).

4. Experimental results

4.1 Performance Comparison to classical ICP

Fig. 7 shows the difference between the results of aligning a hypothetical set of 3 simple scans using classical ICP and our approach. Due to the hard constraints of using the nearest point correspondence only ICP (top row) ends in a non perceptually optimal configuration.

FFS takes into account the correspondences between all points first, a decreasing σ_t finally guarantees the correct positioning of the scans, decreasing the influence of points being too far away. The bottom right image shows the result of FFS after 12 iteration steps.

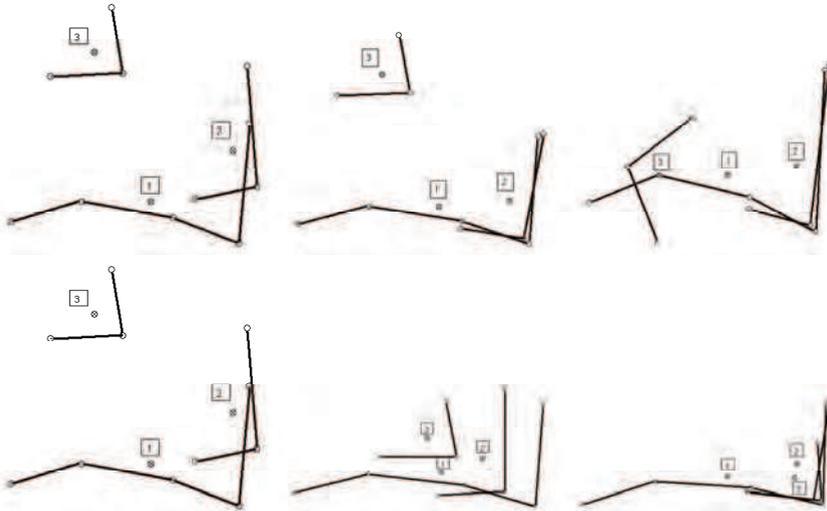


Figure 7. The top row shows 3 steps of the alignment of 3 scans (each scan consists of a single corner only) by classical ICP, the bottom-row shows the results of the proposed approach. The alignment progress can be seen from left to right in both cases. The square boxes show the robot poses of the the scans

4.2 NIST disaster area

The NIST data set used in this experiment simulates a typical data set of multi robot mapping in rescue scenarios. It is especially complicated, as it matches the complicated constraints imposed by these settings, which contain only imprecise odometry, no landmarks and very little overlap.

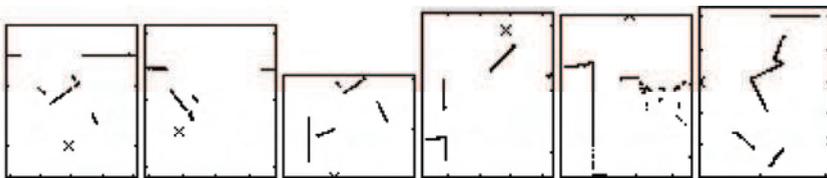


Figure 8. 6 out of 60 scans of the NIST rescue scenario data set. The scans in this data set are very sparse and have minimal overlap

The data set consists of 60 scans taken from 15 different positions in directions E,N,W,S with an overlap of 5° (i.e. overlap between E and N, N and W etc.). The area has a size of approx. $10 \times 15m$, the 15 locations differ approx. $2m$ from each other (see Figure 9). The distance between the positions of the 4 scans taken from an assumed single position differs up to $30cm$, with an angular error of up to 20° to the assumed direction. This data set can be seen as a multi robot mapping scenario using 15 robots, with 4 scans gained from each robot.

Although the pre-alignment assumes this setting, FFS actually treats the 60 scans as independent scans without help of any further information, e.g. constraints on the groups of 4. The single scans have very little pair wise overlap. Figure 9, (l)-(6) shows 6 example scans, all located on the left side of the global map; the overlapping pairs among these scans are (1,2), (1,3),(1,4), (2,3), (3,4), (3,6), (4,5).

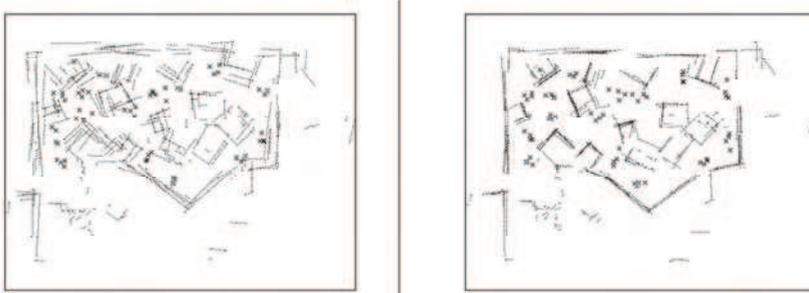


Figure 9. left: 60 scans superimposed building a global map using a rough initial transformation estimation, right: after 20 iterations of FFS. The crosses show the robots' positions, (the reader might try to find the 6 single scans of fig. 8 in the global map. 1,2,3 and 6 are part of the upper left corner, 4 and 5 are located in the lower left corner). The final result of FFS is shown in fig. 10, left

The test performed on this complicated data set demonstrates the robustness of the FFS system. The initial, pre-alignment map is gained by the shape-based algorithm described in Section 3.6. Fig. 9 shows the initial global map as well as iteration step 20, fig. 10 the final global map, after 50 iterations. The data set was re-sampled as described in section 3.3. The radius of the ROI was set to 5 cm, the parameters for the motion cooling were set as:

- Δ_t decreases from 5 to 1 with step factor of 0.96
- σ_t decreases from 15 to 4 with step size of 0.25

Although the data is poorly pre aligned and the overlap between the single scans is minimal, FFS successfully reconstructs the global map, which proves its applicability for this multi robot setting. The mean translation/rotation of the scans (translation/rotation between initial and final global map) is $16\text{cm}/4^\circ$, the maximum translation/rotation is $50\text{cm}/10.5^\circ$.

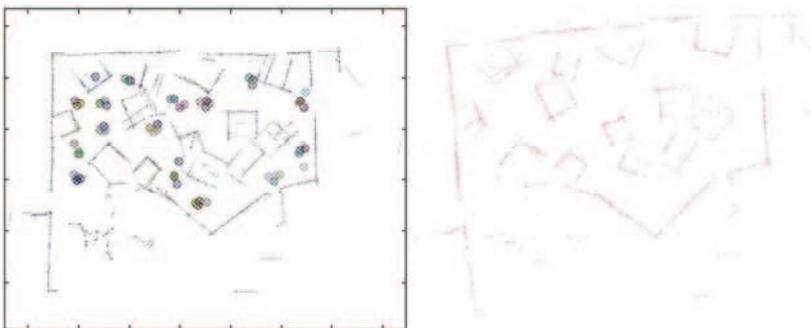


Figure 10. Left, final map (after 50 iterations) of initialization fig. 9 with FFS. Right, the final map obtained by the Lu & Milios technique as reported in (Borrmann et al., 2007). The systems lead to results of comparable quality

The alignment can also be seen as a movie at: <http://knight.cis.temple.edu/~lakaemper/FFS/FFSTheMovie.wmv>

The movie especially makes clear the effect of the motion cooling.

Figure 10 shows the result after 50 iteration steps. The computational time for each iteration is 1 second on a 1.5GHz desktop computer in the current MATLAB implementation, using the bounding box approach as described in section 3.4.1, and resampling with a distance of 10cm.

In order to compare the proposed FFS approach to the state of the art of existing robot mapping approaches, we applied three influential approaches to the NIST data set illustrated in Figure 9. We applied the particle filter based DP-SLAM (Eliazar and Parr, 2004), the ICP based VASCO robot mapping module of CARMEN, and improved grid-based SLAM with Rao-Blackwellized Particle Filters (Grisetti et al., 2005). All three approaches failed to produce any reasonable results, since they are based on sequential processing of data (online SLAM), which can not be applied on this data set due to the extremely minimal overlap of consecutive scans (even if the order is known).

However, we compared to a recent implementation of Lu/Milios type SLAM (Borrmann et al., 2007). The results are shown in figure 10. Both algorithms show an overall comparable performance, although local differences can be seen: the Lu/Milios type SLAM reconstructs the top right corner better, while FFS performs better on the left side.

Figure 12, left, shows the potential $P(g)$ vs. iteration curve for this data set. The potential is monotonically decreasing, hence in this case FFS steers directly towards a (local) minimum, which is reasonable due to the initialization. The next experiment will show a different case.

4.3 Apartment

This experiment demonstrates the benefits and applicability of FFS in data sets which are incorrectly pre aligned e.g. due to effects of wrong loop closing. We used the IROS 2006 test data set taken from <http://staff.science.uva.nl/~zivkovic/FS2HSC/dataset.html>. The data set consists of about 2000 scans from which we select every 10th scan. Thus, our test data set consists of 200 scans taken from a single robot in an apartment of size about 16 x 8m. As shown in Fig. 11(1), the pre-alignment gained shows a huge error, additionally the alignment is very imprecise (blurred features). The experiment shows the power of FFS to escape local minima: starting with a large stepping parameter Δ , the first transformation blurs the data set and therefore weakens the wrong correspondences, giving space for new connections, Fig. 11(2). Transforming all scans in parallel eventually results in a version of the map, which not only shows the misaligned parallel walls correctly contracted but also corrected the huge error as shown in Figure 11(4). The values of the parameters are equal to the experiment in Section 4.2. Figure 11 also shows a limit of the algorithm: one single initially strongly misaligned scan does not find consistent correspondences and therefore can not be correctly re positioned by the algorithm. It stays in its incorrect position. We assume that no algorithm working only on low level perceptual features is able to handle such a strong error correctly; mid level cognitive correspondences are needed. However, mid level perceptual features can easily be integrated into the system using correspondence functions modeling these perceptual forces, which will be part of the future work on the system.

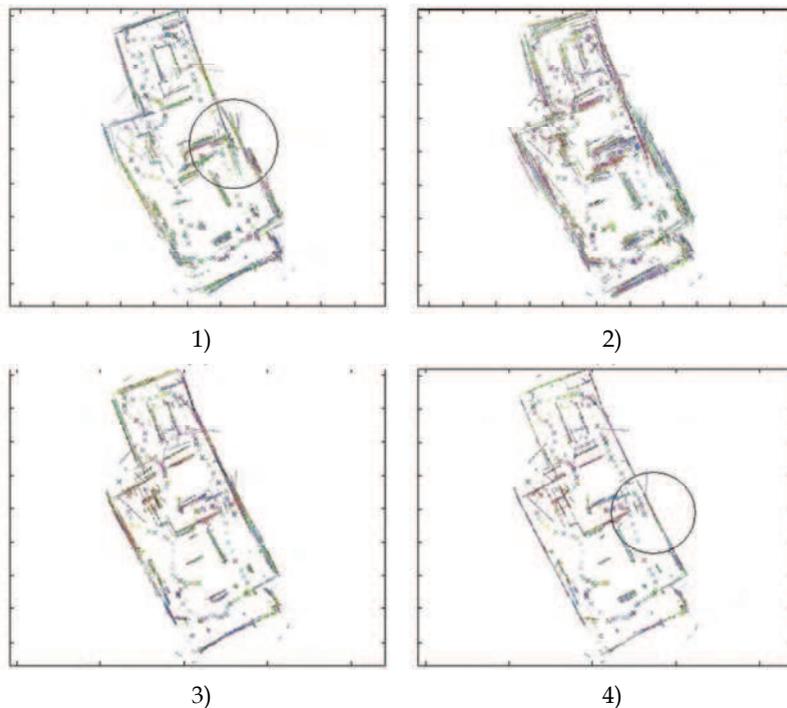


Figure 11. Apartment data set. 1) initial configuration. The circled area shows an error due to incorrect loop closing. 2) A large step parameter dT blurs the map in the first iteration step to escape from the local minimum 3) iteration 50: 4) iteration 150: FFS has not only contracted the edges given in 1), but also has realigned the entire global map to fix the error (circled area)

4.4 NIST maze

This data set consists of 16 scans with similar structures, a typical indoor environment, yet again scanned with minimal overlap. See figure 13 and 14 for this experiment.

5. Conclusion and future work

We presented a new approach to the problem of multi robot mapping under the constraints given in rescue scenarios. It does not rely on odometry i.e. the relative pose between the robots is unknown. It also can deal with the problem of extremely minimal overlap. Experiments conducted on a real data set of a disaster area from NIST shows the performance of the FFS approach under these complicated constraints and proved its applicability to the problem of multi robot mapping, they also proved the excellent performance of the algorithm correcting effects from wrong pre alignment. The future work will mainly focus on detection of higher level features: the modeling of the correspondence function with respect to the masses opens different ways to interface to mid level modules. The approach is easily extendable to 3D, a report about the performance of an implementation of the 3D FFS is the topic of a forthcoming paper.

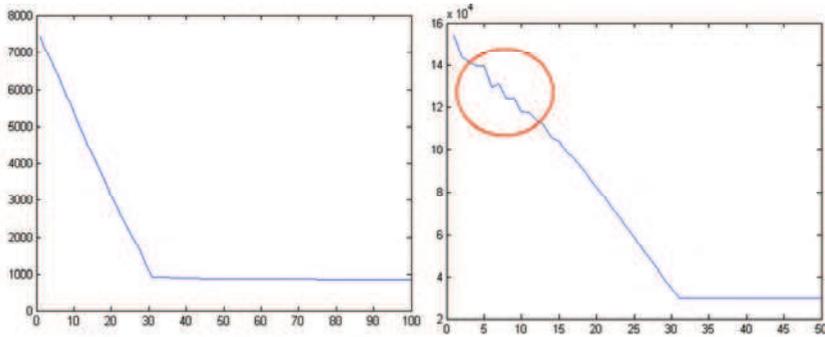


Figure 12: Left, potential vs. iterations of FFS for disaster data. Right, potential for Apartment data set. The potential (encircled) of the apartment data is not monotonically decreasing, indicating a possible escape from a local minimum

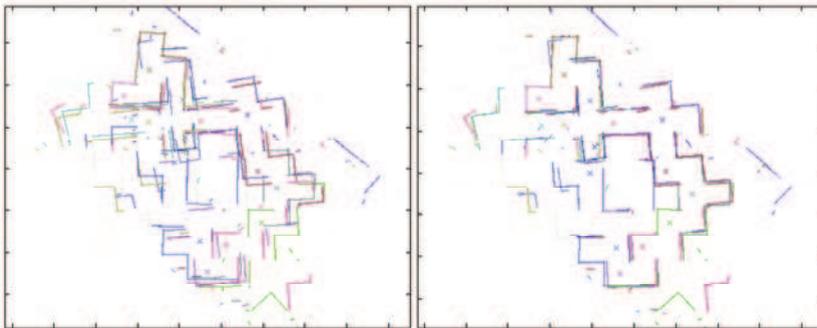


Figure 13. Left, initial configuration of NIST's maze data (16 scans). Right, after 5 iterations of FFS

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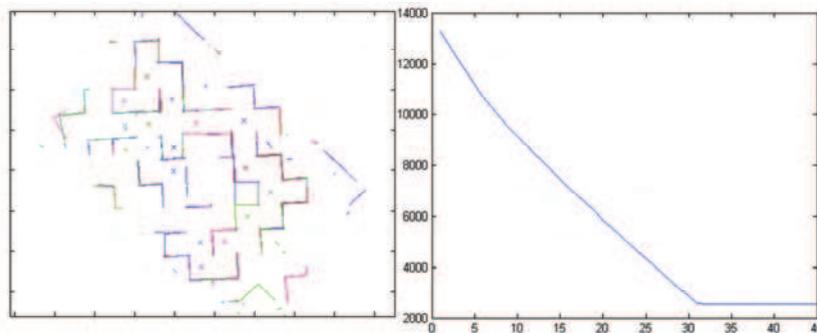


Figure 14. Left, final map obtained with FFS. Right, the potential vs. iterations plot

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To design a team of robots which is able to perform given tasks is a great concern of many members of robotics community. There are many problems left to be solved in order to have the fully functional robot team. Robotics community is trying hard to solve such problems (navigation, task allocation, communication, adaptation, control, ...). This book represents the contributions of the top researchers in this field and will serve as a valuable tool for professionals in this interdisciplinary field. It is focused on the challenging issues of team architectures, vehicle learning and adaptation, heterogeneous group control and cooperation, task selection, dynamic autonomy, mixed initiative, and human and robot team interaction. The book consists of 16 chapters introducing both basic research and advanced developments. Topics covered include kinematics, dynamic analysis, accuracy, optimization design, modelling, simulation and control of multi robot systems.

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