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The variational counterpart of the equation above is

$$a(u, v) + \lambda m(u, v) = f(v)$$

for all $v \in H$, where

$$a(u, v) := \langle \mathcal{G}u, v \rangle_{H' \times H} \text{ and } m(u, v) := \langle u, v \rangle_{H \times H} .$$

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In typical situations, the bilinear form $a(\cdot, \cdot)$ representing the integral operator can be written as

$$a(u, v) = \int_{\Omega} v(x) \int_{\Omega} g(x, y) u(y) dy dx \quad (3)$$

for a kernel function $g(\cdot, \cdot)$ and a domain or manifold Ω .

The equation (2) is discretized by choosing an n -dimensional subspace H_n of H and considering the problem of finding a function $u_n \in H_n$ s.t.

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For any basis $(\varphi_i)_{i \in \mathcal{I}}$ of H_n , this is equivalent to

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Since the solution u_n is an element of H_n , there is a coefficient vector $(x_i)_{i \in \mathcal{I}}$ satisfying

$$u_n = \sum_{j \in \mathcal{I}} x_j \varphi_j,$$

s.t. the coefficients satisfy the equation

$$\sum_{j \in \mathcal{I}} x_j a(\varphi_j, \varphi_i) + \lambda \sum_{j \in \mathcal{I}} x_j m(\varphi_j, \varphi_i) = f(\varphi_i)$$

for all $i \in \mathcal{I}$.

This is a system of linear equations and can be written in matrix form

$$Gx + \lambda Mx = b$$

by introducing matrices $G, M \in \mathbb{R}^{\mathcal{I} \times \mathcal{I}}$ and a vector $b \in \mathbb{R}^{\mathcal{I}}$ with

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Storing G directly doesn't lead to efficient algorithms. Therefore we approximate G by a matrix that can be treated efficiently, by replacing the kernel function $k(\cdot, \cdot)$ by local degenerate approximations, and this leads to a hierarchical matrix.

Recall: Degenerate approximation

Last week we've already talked about degenerate approximation, but as I can imagine that a few people in here may not remember it very vividly, I'll quickly explain what it is about.

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The idea is to approximate the kernel function $g(\cdot, \cdot)$ by using interpolation instead of Taylor expansion, and thus avoiding the need of being able to evaluate the derivatives of g efficiently.

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$(x_\nu)_{\nu \in K}$ be a family of interpolation points in \mathbb{R}^d

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We define

$$\tilde{g}(x, y) := \sum_{\nu \in K} g(x_\nu, y) \mathcal{L}_\nu(x).$$

As mentioned before, we don't need any derivative of g for that approximation.

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We can now replace the matrix G with \tilde{G} defined by

$$\tilde{G}_{ij} := \int_{\Omega} \varphi_i(x) \int_{\Omega} \tilde{g}(x, y) \varphi_j(y) dy dx = (AB^T)_{ij},$$

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Part II: Boundary Element Method in 2D

We consider a closed curve in 2-dimensional space, given as an array vertex of n points. We use piecewise constant basis functions and choose the characterizing point to be the middle of the corresponding interval. We will now solve integral equations on this curve.

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We are interested in a boundary integral problem, i.e. the set Ω is a submanifold. Here Ω is a curve. Since we'll be talking about integral equations, I will recall the meaning of an integral on a curve:

1 Curve integrals

Let $\gamma : [0, 1] \rightarrow \mathbb{R}^2$ be injective in $[0, 1[$, $\gamma \in C^1, \gamma' \in C^0$. We write $\Gamma := \gamma([0, 1])$. Let $u \in C^0(\Gamma)$. We introduce a partition $0 = x_0 < x_1 < \dots < x_n = 1$ of $[0, 1]$ and consider the sum

$$I_x := \sum_{i=1}^n u(\gamma(x_i)) \|\gamma(x_i) - \gamma(x_{i-1})\|.$$

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Lemma (Curve integral) : *Let $\epsilon \in \mathbb{R}_{>0}$. There is a $\delta \in \mathbb{R}_{>0}$ s.t. \forall partitions $0 = x_0 < x_1 < \dots < x_n = 1$ with $x_i - x_{i-1} < \delta$ ($i \in 1, \dots, n$) we have*

$$\|I_x - \int_0^1 u(\gamma(y)) \|\gamma'(y)\| dy\| \leq \epsilon.$$

Proof: elementar analysis

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Proof: elementar analysis

Definition: We define the *curve integral* : let $(\gamma_i)_{i=1}^m$ be a tuple of injective functions in $C^1([0, 1], \mathbb{R}^2)$. For all $i \in 1, \dots, m$, we set $\Gamma_i := \gamma_i([0, 1])$. The *curve integral* over the piecewise differentiable curve $\Gamma := \cup_{i=1}^m \Gamma_i$ is given by

$$\int_{\Gamma} u(x) dx := \sum_{i=1}^m \int_0^1 u(\gamma_i(y)) \|\gamma_i'(y)\| dy.$$

2 Single layer potential

We fix n points $p_0, \dots, p_{n-1} \in \mathbb{R}^2$, set $p_n := p_0$ and define the affine parametrizations

$$\gamma_i : [0, 1] \rightarrow \mathbb{R}^2, \quad y \mapsto p_{i-1}(1 - y) + p_i y,$$

for $i \in 1, \dots, n$. As long as $p_i \neq p_j$ holds $\forall i, j \in 0, \dots, n - 1$ with $i \neq j$, this defines a polygonal curve $\Gamma := \cup_{i=1}^n \gamma_i([0, 1])$.

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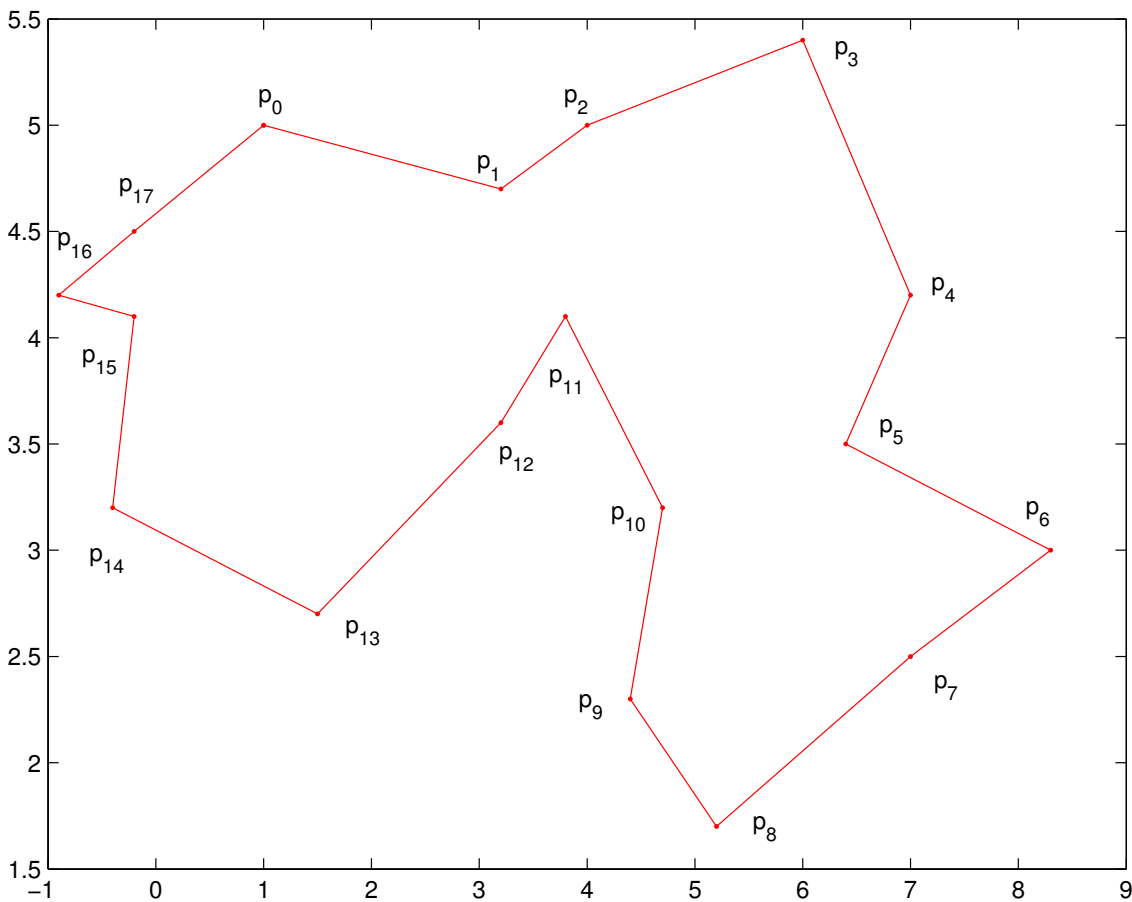
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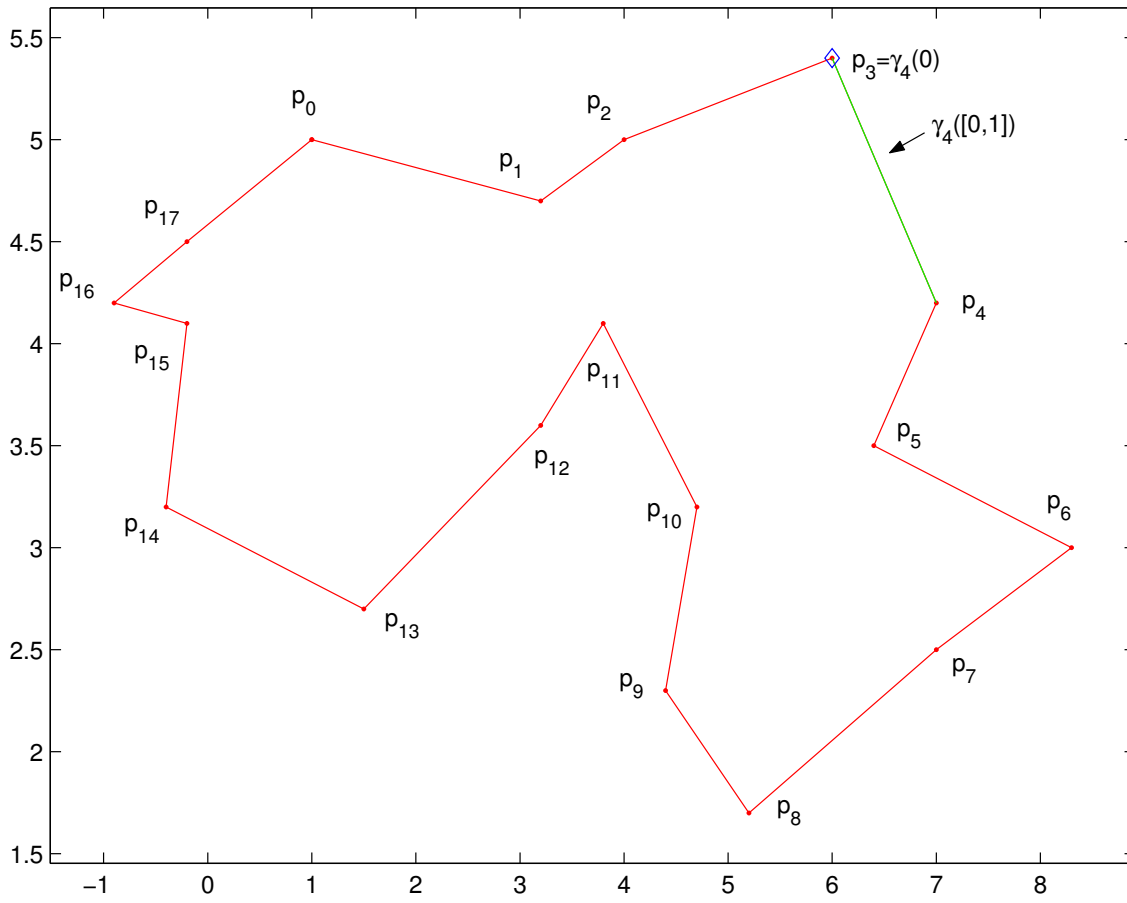
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Example:

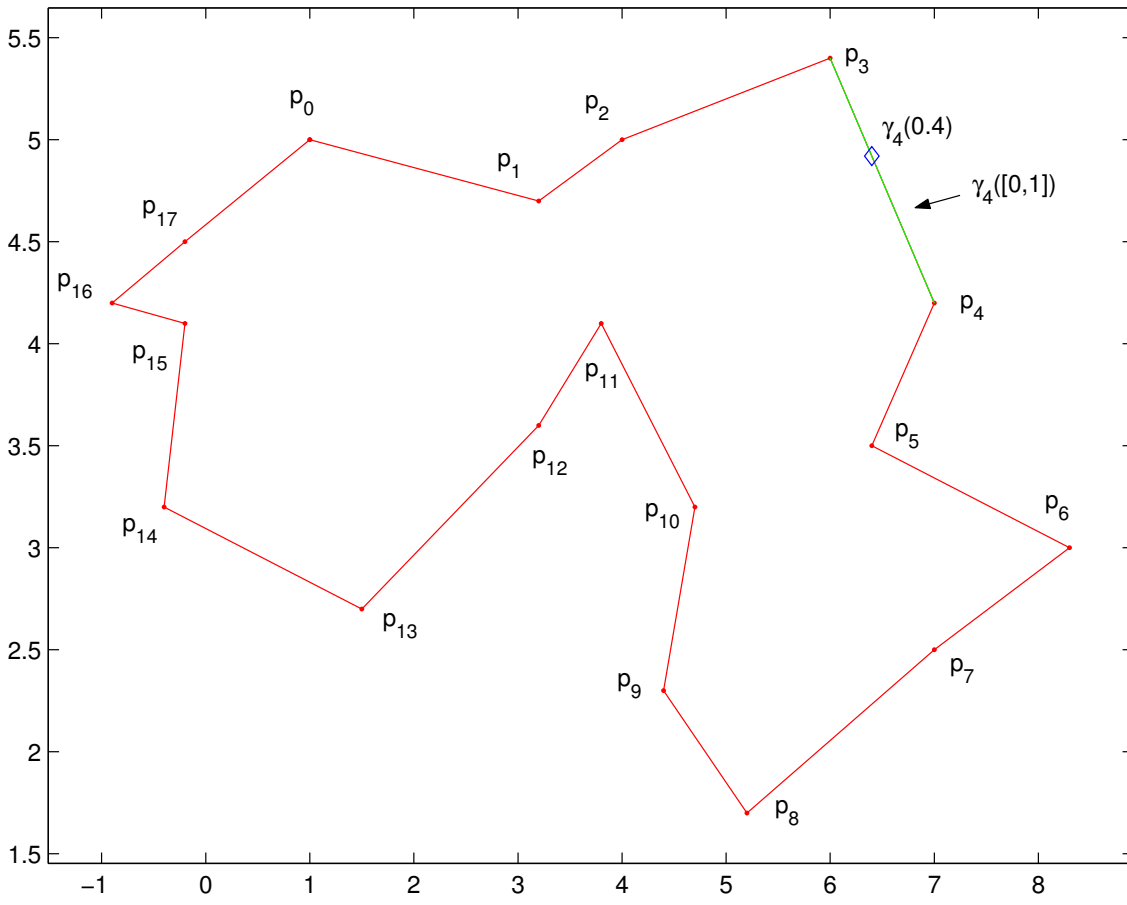
We consider the following points:



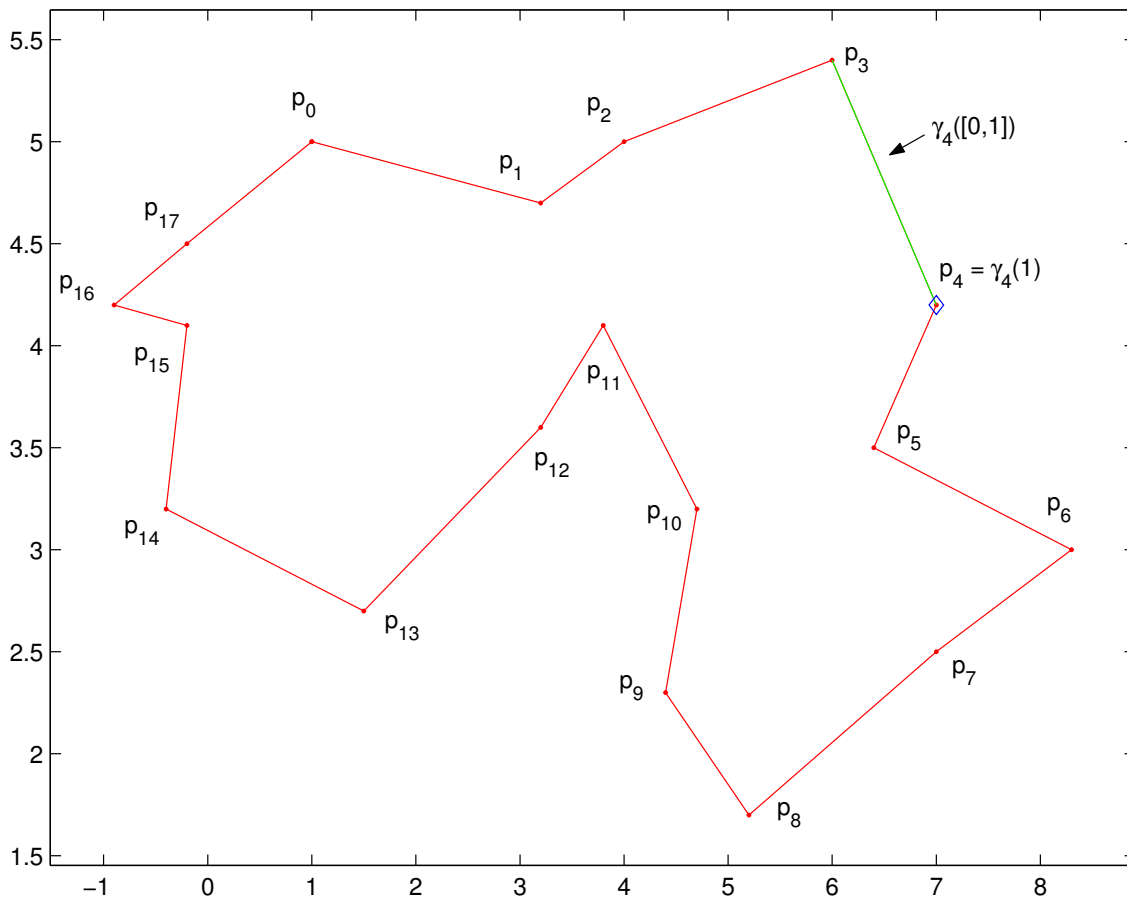
We observe γ_4 and its evaluation in $t=0$:



$t=0.4:$



and $t=1$:



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We discretize $a_{slp}(\cdot, \cdot)$ using piecewise C^{∞} functions $(\varphi_i)_{i=1}^n$ defined through

$$\varphi_i \circ \gamma_j \equiv \delta_{ij}$$

for $i, j \in \mathcal{I} := 1, \dots, n$. The coeff. of the corresponding matrix are given by

$$\begin{aligned} G_{ij} &= a_{slp}(\varphi_i, \varphi_j) = \int_{\Gamma} \varphi_i(x) \int_{\Gamma} \log(\|x - y\|)\varphi_j(y)dydx \\ &= \|p_i - p_{i-1}\| \|p_j - p_{j-1}\| \int_0^1 \int_0^1 \log(\|\gamma_i(x) - \gamma_j(y)\|)dydx. \end{aligned}$$

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From $p_i \neq p_j$ it follows that this matrix is full. As long as $\gamma_i([0, 1])$ doesn't intersect $\gamma_j([0, 1]) \forall i, j \in 1, \dots, n$, we don't have singularities and can replace the kernel by degenerate approximations.

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They correspond to admissible pairs (t,s) of clusters and require the evaluation of a degenerate approximation of the kernel function. Let's assume that $\text{diam}(Q^t) \leq \text{diam}(Q^s)$, it follows

$$\tilde{g}(x, y) = \sum_{\nu \in K} \log(\|x_\nu^t - y\|) (L)_\nu^t(x)$$

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γ_i is affine, so $A_{i\nu}^{t,s}$ are polynomials of degree m . We can thus use an exact quadrature rule for its evaluation.

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- An array `l` of dim. `p` containing the transformed interpolation pts.

Recall(Gauss-quadrature):

The idea is to approximate an integral as

$$\int_{\Omega} f(\xi) d\xi \approx \sum_{K \in \mathcal{M}} |K| \sum_{l=1}^{P_K} w_l^K f(\pi_l^K).$$

The w_l^K are called local weights and the points π_l^K local nodes. The Gauss-quadrature is exact for polynomials of degree up to $2P - 1$ using only P nodes.

The quadrature is numerically stable if all the weights are positive.

Example(Gauss-quadrature, 1-dimensional):

We consider the integral $\int_{-1}^1 f(x)dx$ and want to approximate it with a Gauss-quadrature of degree $P=2$:

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$$\int_{-1}^1 f(\xi)d\xi = w_1f(x_1) + w_2f(x_2)$$

This approximation has to be exact for polynomials $\in \mathcal{P}_3$, so we choose successively $f(x) = x^0, x^1, x^2, x^3$. We now have 4 equations to solve:

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And so we have

$$\begin{aligned} w_1 &= w_2 = 1 \\ x_1 &= -\frac{1}{\sqrt{3}} \\ x_2 &= \frac{1}{\sqrt{3}} \end{aligned}$$

The evaluation of $B^{t,s}$ requires us to integrate the kernel function for points x_ν^t on intervals given by p_{i-1} and p_i . Here, it can be done analytically. In more general cases, we can use the same quadrature rule as for polynomials, but the result will no longer necessarily be exact.

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Example(affine γ in \mathbb{R}^2): Let $\gamma : [0, 1] \rightarrow \mathbb{R}^2$ and $c \in \mathbb{R}^2$ given by

$$\gamma(t) := \begin{pmatrix} s_x + td_x \\ s_y + td_y \end{pmatrix} \text{ and } c := \begin{pmatrix} c_x \\ c_y \end{pmatrix}.$$

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and if we change γ to

$$\gamma(t) := \begin{pmatrix} 1 + 2t \\ 1 + 4t \end{pmatrix}$$

(and thus doubling its"speed") we obtain $b=5.4231$.

The `supermatrix` structure can be initialized by a simple recursion: If the `supermatrix` contains an `rkmatrix`, we compare the diameters of the clusters involved and use the procedure described above to initialize the fields `a` and `b` of the `rkmatrix`.

If the `supermatrix` contains a `fullmatrix`, we evaluate singular integrals and fill its field `e`.

Otherwise, we proceed recursively with the subblocks that are given by the array `s`.