

Chloride diffusivity in hardened cement paste from microscale analyses and accounting for binding effects

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Abstract. The diffusion of chloride ions in hardened cement paste (HCP) under steady-state conditions and accounting for the highly heterogeneous nature of the material is investigated. The HCP microstructures are obtained through segmentation of X-ray images of real samples as well as from simulations using the cement hydration model CEMHYD3D. Moreover, the physical and chemical interactions between chloride ions and HCP phases (binding), along with their effects on the diffusive process, are explicitly taken into account. The homogenized diffusivity of the HCP is then derived through a least square homogenization technique. Comparisons between numerical results and experimental data from the literature are presented.

Keywords: Chloride diffusion and binding, Microscale analyses, Numerical analyses, Segmentation of real microstructures, Simulated microstructures

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1. Introduction

The diffusion of chlorides in reinforced concrete structures often controls their durability and its prediction is needed both to forecast service life and to schedule an efficient maintenance strategy [1–3]. Chloride ions can be found in hardened cement paste (HCP) either because they are present in the ingredients (water, cement clinker, supplementary cementitious materials, and/or aggregates), or as a consequence of diffusion from the external environment [4–6]. Other than using a chloride-based accelerator, the former case is mostly related to a modification of the production process of the cement clinker,

9 but is not likely to occur in typical practice since the presence of chlorides in concrete
10 raw materials is generally restricted by production guidelines [4, 6]. Conversely, the
11 penetration of chlorides from the external environment is of more practical interest
12 since it is representative of many aggressive environments such as exposure to de-icing
13 salts or marine/tidal environments [4].

14 The penetration of chlorides from the external environment is strongly influenced
15 by the characteristics and the heterogeneous nature of the HCP at the microscopic
16 level [7–9] (e.g., at a length scale of 1 μm and larger). In particular, the topological
17 and physical characteristics of the capillary pore system [7, 10] and the possibility for
18 the ions to have easy access to the pores are among the most important parameters
19 governing the diffusion rate [7]. Also, the solid phases play a significant role for the
20 chloride penetration into HCP, as the microporosity of the solid matrix constitutes a
21 slower but effective pathway for ionic diffusion. Their effect becomes more important as
22 the capillary porosity decreases, reaching a primary role in the HCP where the capillary
23 pore system does not form a continuous (percolated) pathway in three dimensions [7].
24 Moreover, it is well known that some products of the hydration of the cement clinker can
25 physically or chemically interact with chloride ions, modifying the diffusion processes
26 through the so-called binding effect [1, 3, 4, 6, 11, 12].

27 The level of detail needed to study the diffusion at the microscopic level adds
28 considerable complexity to experimental tests and their interpretation. For this reason,
29 the problem is often addressed by computational modeling. The diffusion problem
30 is classically addressed by adopting Ficks theory under some simplifying assumptions
31 such as one-dimensional flow or the absence of binding (see [12–17] among others).
32 More recently, Garboczi and Bentz [8] proposed a numerical model for diffusion based
33 on the electrical conductivity-diffusivity analogy and discretization of the HCP using a
34 3-D lattice of conductors, each with a different diffusivity/conductivity as a function of
35 the underlying phase. Based on such a model, an empirical relationship was developed
36 to evaluate the diffusivity of the HCP at the mesoscale as a function of the capillary
37 porosity. The same authors present in [1] a different model based on random walk
38 theory applied to a discretized two-dimensional microstructure that also accounts for the
39 interaction between chlorides and the solid phases. The electrical conductivity analogy
40 and random walk approaches were subsequently combined [18] in a multiscale approach
41 where the diffusivity of bulk cement paste and of the interfacial transition zone (ITZ)
42 between the paste and the aggregates are defined at the microscale through the formula
43 presented in [1], while accounting for the aggregate volume fraction at the mesoscale.
44 Then, random walk theory similar to that proposed in [1] is used to simulate diffusion
45 at the mesoscale, which is conceived as a three phase continuum medium composed of
46 non-diffusive aggregates, bulk cement paste, and the ITZ. The model proposed in [18]
47 is subsequently simplified through the introduction of analytical procedures [19].

48 A multiscale method is also adopted by Zhang et al. [9, 20], where the authors
49 use Lattice-Boltzmann and finite element methods respectively for the micro- and
50 mesoscales. Here, the authors use simulated microstructures and account also for the

diffusive behavior of the ITZ, however the binding effect is treated in a very simplified manner. Nilenius et al [21, 22] study the upscaling of the diffusive properties from the meso- to the macroscale, neglecting the microscale and thus the binding effect. The interaction between diffusion, binding, and the presence of cracks is investigated for a 3D structure broadly segmented into HCP and voids by Lu et al. in [23]. Finally, Ukrainczyk and Koenders in [10] investigate the mass transport at the microscale level and the role of the capillary pore system topology neglecting binding, while in [24] they employ a similar approach to study the transient phase of the calcium ion leaching process in HCP.

The main aim of the present work is to study the steady-state diffusion behavior of externally supplied chlorides in HCP accounting for its highly heterogeneous microstructure, as well as for physical and chemical binding effects acting at the single phase level (i.e., considering the various chemical compounds present). To this end, Ficks law of diffusion is modified through the consistent introduction of a phase-dependent binding activity coefficient. The 3D HCP microstructures for the analyses are both simulated, using CEMHYD3D [25, 26], as well as segmented starting from real X-ray computer tomography (CT) images available from the NIST Visible Cement Database [27, 28]. The model is implemented in a finite difference code and several tests are simulated varying the environmental chloride concentration. The diffusivity at the mesoscale is evaluated through a linear homogenization using both experimental data and widely accepted analytical relationships from the literature. A comparison between results from real and simulated HCP microstructures is also performed and discussed.

The paper is structured as follows: in section 2 the governing equations of diffusion coupled with binding are presented. In section 3 the effects of the binding are described and quantitatively evaluated. The numerical aspects of the implementation and the simulations performed are described respectively in section 4 and 5. Finally, some concluding remarks are made in section 6.

2. Chloride diffusion and binding

2.1. Governing equations for chloride diffusion in HCP

The migration of particles inside a medium should pointwise satisfy the general mass conservation condition, which reads

$$\frac{\partial C}{\partial t} = -\nabla \cdot \mathbf{J}, \quad (1)$$

where C is the particle concentration, t is the time and \mathbf{J} is the flux vector.

To completely describe the motion of the particles, a flux constitutive relation is needed. When dealing with diffusion of ions in porous media, such as for the case of chloride diffusion in HCP studied here, the flux constitutive equation can be easily formulated taking into account the conditions in which the migration of ions takes place. Firstly, water saturated conditions are assumed since they are imposed in most

89 experimental setups and occur in many real-world cases due to the limited thickness
 90 of the concrete cover. Moreover, in absence of external forces inducing the motion of
 91 the fluid, advection can be neglected, leading to a pure diffusive process. It is also
 92 assumed here that the principal driving force responsible for the diffusion process is the
 93 ions concentration gradient, meaning that all other potential fields (e.g., electrical fields,
 94 presence of temperature or chemical potential gradients) are negligible.

95 With the above assumptions, the constitutive flux equation can be expressed
 96 through Fick's first law as

$$97 \quad \mathbf{J} = -D\nabla C, \quad (2)$$

98 where D is the diffusion coefficient or diffusivity.

99 The definition of the diffusivity parameter D in Equation 2 is strongly related to
 100 the problem at hand. In particular, it is possible to distinguish between

- 101 - D^* or self-diffusion coefficient: it is defined as the diffusion coefficient of a species
 102 when the chemical potential gradient equals zero. It describes the mobility of a
 103 particle within other particles of the same species, i.e. it is a macroscopic description
 104 of the Brownian motion (bulk motion in homogeneous media);
- 105 - D_{free} or free diffusivity (also, inter-diffusivity): it describes the diffusivity of a
 106 binary system composed of the diffusive species and a solvent. It is related to the
 107 self-diffusion coefficient by the equation

$$108 \quad D^* = D_{free} \frac{\partial \ln C_{diff}}{\partial \ln a_{diff}}, \quad (3)$$

109 where C_{diff} and a_{diff} are the concentration and the chemical activity of the diffusive
 110 species. What is usually taken as a reference value is the diffusivity of a species
 111 into pure water at a constant temperature;

- 112 - D_c or diffusivity in saturated capillary pores: when migration takes place into small
 113 pores (i.e., diameter $\leq 5 \mu\text{m}$ to $10 \mu\text{m}$), phenomena such as particle-environment
 114 interaction and the viscosity of the fluid become non negligible, thus reducing the
 115 diffusivity with respect to D_{free} ;
- 116 - D_s or solid diffusivity: diffusivity can take place also in those solid materials
 117 where the inner structural arrangement (e.g., clusters of crystal grains) involves the
 118 presence of micro- or nano-capillaries whose geometry cannot be resolved explicitly.
 119 In such cases, the value of diffusivity is defined by decreasing the value of D_c to
 120 account for the topology, tortuosity and volume fraction of the capillary system in
 121 the solid domain.

122 The choice of the correct diffusivity parameter to be used is mainly related to the
 123 observation scale and the nature of the solvent, the diffusive species and the media
 124 hosting the diffusion. However, since the validity of Equations 1 and 2 is general, in the
 125 following mathematical formulation the diffusivity coefficient will be generally termed
 126 D and its meaning and definition will be specified in the validation section according to
 127 the problem studied.

128 Substituting Equation 2 into Equation 1 gives

$$129 \quad \frac{\partial C}{\partial t} = \nabla \cdot D \nabla C, \quad (4)$$

130 which is termed Fick's second law. In homogeneous materials, D is usually assumed
 131 to be a constant, i.e. independent from position, concentration and time. However, as
 132 already introduced in Section 1, HCP is composed of different hydrated and unhydrated
 133 phases, each characterized by a very different diffusion behavior, hence

$$134 \quad D = D(\mathbf{x}), \quad (5)$$

135 where \mathbf{x} is the spatial coordinate vector.

136 2.2. Governing equations in presence of binding

137 Equation 4 accounts for the heterogeneous nature of the HCP, but neglects any possible
 138 chemical and physical interaction between the diffusive species and the phases in which
 139 diffusion takes place. In particular, it has been extensively demonstrated (see e.g.
 140 [1, 4, 11, 29]) that chloride ions penetrating into the HCP from the environment can
 141 interact with some hydration products, thereby modifying the diffusion process. In
 142 particular, the total concentration of chloride ions C_{tot} can be written as the sum of two
 143 terms

$$144 \quad C_{tot} = C_f + C_b, \quad (6)$$

145 where C_f and C_b are the concentrations respectively of free (i.e., the ions that are free to
 146 migrate) and bound chlorides. The latter are permanently or dynamically bound to the
 147 solid phase, thus they cannot take part in the diffusion process anymore because they
 148 have become part of the solid mass of the HCP. On the other hand, the mass balance
 149 should account for the total concentration of particles C_{tot} . Hence, Equation 4 can be
 150 rewritten as

$$151 \quad \frac{\partial C_{tot}}{\partial t} = \nabla \cdot D(\mathbf{x}) \nabla C_f. \quad (7)$$

152 Considering then the relationship

$$153 \quad \nabla C_f = \nabla(C_{tot} - C_b) = \left(1 - \frac{\partial C_b}{\partial C_{tot}}\right) \nabla C_{tot}, \quad (8)$$

154 Equation 7 becomes

$$155 \quad \frac{\partial C_{tot}}{\partial t} = \nabla \cdot \left[\left(1 - \frac{\partial C_b}{\partial C_{tot}}\right) D(\mathbf{x}) \nabla C_{tot} \right]. \quad (9)$$

156 It is now possible to introduce the effective diffusivity D_{eff} as

$$157 \quad D_{eff}(\mathbf{x}, C_{tot}) = \left(1 - \frac{\partial C_b}{\partial C_{tot}}\right) D(\mathbf{x}) = \mu_B(C_{tot}) D(\mathbf{x}), \quad (10)$$

158 where

$$159 \quad \mu_B(C_{tot}) = 1 - \frac{\partial C_b}{\partial C_{tot}}, \quad (11)$$

160 is the binding activity coefficient, which makes the diffusivity coefficient dependent on
 161 the capacity of the phase hosting the diffusion to bind the diffusive species. However,
 162 as will be better explained later on, the amount of chlorides bound by the solid phase in
 163 HCP is generally expressed as a function of C_f instead of C_{tot} , i.e. $C_b = C_b(C_f)$. As a
 164 result, the binding activity coefficient as defined in Equation 11 is not readily available
 165 and, hence, a further manipulation is needed

$$166 \quad \mu_B = 1 - \frac{\partial C_b}{\partial C_{tot}} = \frac{\partial C_f}{\partial C_{tot}} = \left[\frac{\partial (C_f + C_b)}{\partial C_f} \right]^{-1} = \left(1 + \frac{\partial C_b}{\partial C_f} \right)^{-1}. \quad (12)$$

167 Introducing Equation 10 into Equation 9 and accounting for Equation 12 yields

$$168 \quad \frac{\partial C_{tot}}{\partial t} = \nabla \cdot D_{eff}(\mathbf{x}, C_f) \nabla C_{tot}, \quad (13)$$

169 that should be coupled with

$$170 \quad C_f + C_b(C_f) = C_{tot}. \quad (14)$$

171 The introduction of the above non-linear equation (Equation 14) is needed since the
 172 differential field equation (Equation 13) is solved for the unknown C_{tot} , while the
 173 effective diffusivity $D_{eff}(\mathbf{x}, C_f)$ is a function of the free chloride concentration C_f . Once
 174 Equation 14 is solved, the binding activity coefficient $\mu_B(C_f)$ of Equation 12 can be
 175 updated to solve again Equation 13. Both Equation 13 and 14 are here solved using a
 176 Newton-Raphson algorithm.

177 2.3. Steady-state conditions

178 The diffusion process is composed of two stages: an initial transient phase where the
 179 flux of the diffusive species is established and the concentration changes in time, and a
 180 steady-state phase, where a steady flux is achieved and the variation over time of the
 181 concentration is zero.

182 Since the practical interest in studying chloride diffusion and chloride binding in
 183 HCP is mostly related to long-term effects (e.g., corrosion of the steel rebars and other
 184 slowly evolving phenomena), in the following steady-state conditions are assumed. In
 185 particular, the effects of the interaction between cement paste phases and chlorides on
 186 the effective diffusivity are investigated.

187 Steady-state conditions are enforced imposing that the time derivative in
 188 Equation 13 vanishes, thus obtaining

$$189 \quad \nabla \cdot D_{eff} \nabla C_{tot} = \nabla \cdot \mu_B D \nabla C_{tot} = 0. \quad (15)$$

190 3. Binding mechanisms and related equations

191 Although other minor mechanisms are reported in the literature [4, 6, 30, 31], the
 192 binding of chloride ions by the hydration products in HCP follows two main mechanisms,
 193 i.e. chemical and physical interaction. Binding mechanisms, whose development is
 194 influenced by different factors [6, 32, 33], will be illustrated in detail for each phase
 195 involved in section 3.1.

196 The chemo-physical processes responsible for the chloride uptake are not
 197 instantaneous, but develop in time at a certain speed called binding rate [2, 11]. This
 198 rate depends on the environmental conditions and on the interacting compounds. Some
 199 authors tried to empirically calibrate the binding kinetics for the whole HCP through
 200 linear relationships, using results from diffusion tests [2]. Nevertheless, the literature
 201 available to date does not provide reliable data on the binding rate. However, it can be
 202 assumed that the binding process is fast compared to the transient diffusion phase [2],
 203 i.e. that the time needed for its full development is significantly less than that required
 204 for the establishment of the steady-state regime. Hence, it is reasonable to assume that
 205 the maximum amount of chloride ions bound for a given phase has already been reached
 206 once steady-state conditions are attained.

207 As reported by many authors (e.g. [6, 11, 32]), among all the hydration products
 208 present in ordinary Portland cement (OPC), only two are (almost entirely) responsible
 209 for the chloride uptake in HCP: the *CSH* (calcium silicate hydrates) gel and the
 210 *AFm* species (e.g., monosulfate hydrate), while the contributions of *Aft* phases (e.g.,
 211 ettringite), *CH* (portlandite) and other hydrated or unhydrated phases are negligible
 212 under most conditions.

213 *CSH* gel is produced by the hydration of silicates (C_3S and C_2S) and is responsible
 214 for the physical binding of chloride ions [26, 34]. Conversely, the *AFm* presence is
 215 regulated by the amount of aluminates (C_3A and C_4AF) in the clinker, as well as the
 216 contents and forms of sulfates and carbonates [34].

217 3.1. Main binding mechanisms involved

218 Physical binding or adsorption is due to the mutual attraction between charged particles
 219 (i.e., Van der Waals forces) and is strongly related to the extent of the free *CSH* surface
 220 area.

221 Chemical binding takes place by chemical reaction [11, 32], with the presence
 222 of chloride ions leading to the formation chloroaluminate phases in preference to the
 223 conventionally formed sulfoaluminate ones structure (e.g., *AFm* products as $C_3A \cdot$
 224 $CaSO_4 \cdot 12H_2O$). For example, the chemical reaction:



226 that produces Friedel's salt ($C_3A \cdot CaCl_2 \cdot 10H_2O$). According to Equation 16, one mole
 227 of monosulphate is able to bind 2 moles of chlorides [11], but this is a theoretical result
 228 that is reached only asymptotically because it is abstracted from the real environment

Table 1. Specific and molar masses used to convert the binding isotherms.

	Chloride Cl^-	AFm	CSH	HCP
γ [kg/m^3]	3.21	1990.00	2120.00	1800.00
m [g/mol]	35.45	622.87	228.96	-

where the reaction takes place. Chemical binding is stronger than physical binding, since it is due to chemical interactions instead of polar attraction, and is more likely to be irreversible.

3.2. Binding isotherms

The relationship between free and bound chlorides $C_b(C_f)$ at a given temperature for a defined material (single phase or HCP) is termed a binding isotherm. In the present work the isotherms proposed by Hirao et al. [11] are assumed to hold and they are converted to $\mu g/mm^3$ using the molar masses m and the specific masses γ listed in Table 1. The isotherms adopted are depicted in Figure 1a, while the $\mu_B(C_f)$ curve is shown in Figure 1b.

3.2.1. CSH gel For the CSH gel the following isotherm was proven to adequately fit the experimental results [11] (Figure 1a)

$$C_b^{CSH} = K_{CSH} \frac{a_{CSH} C_f}{1 + a_{CSH} C_f}, \quad (17)$$

where concentrations are expressed in $\mu g/mm^3$ and $K_{CSH} = 46.30$ and $a_{CSH} = 7.47 \cdot 10^{-2}$ are two experimentally calibrated parameters [11]. Also, following the definition of binding activity coefficient in Equation 11, the following expression is obtained for the CSH (Figure 1b)

$$\mu_{CSH} = \left[1 + \frac{K_{CSH} a_{CSH}}{(1 + a_{CSH} C_f)^2} \right]^{-1}. \quad (18)$$

3.2.2. AFm phase For the AFm phase a power-law isotherm was proposed in [11]

$$C_b^{AFm} = K_{AFm} C_f^\alpha, \quad (19)$$

where concentrations are expressed in $\mu g/mm^3$ and $K_{AFm} = 12.30$ and $\alpha=0.58$ are calibrated experimentally [11]. Furthermore, from Equation 11 the binding activity coefficient is (Figure 1b)

$$\mu_{AFm} = (1 + K_{AFm} \alpha C_f^{\alpha-1})^{-1}. \quad (20)$$

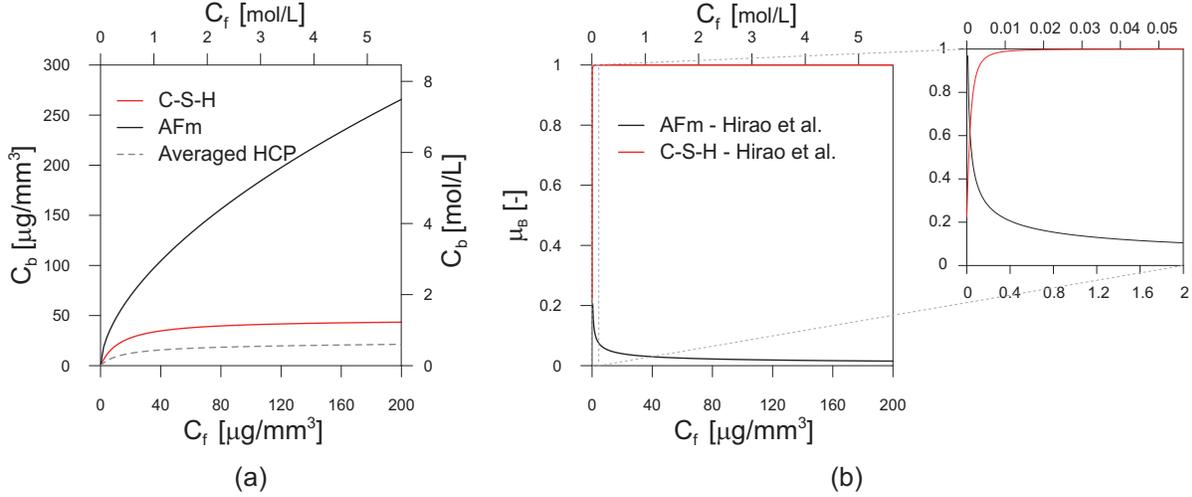


Figure 1. (a) Adopted binding isotherms (from Hirao et al. [11]) and (b) the related binding activity coefficient μ_B .

253 *3.2.3. Averaged HCP isotherm* In their work, Hirao et al. [11], proposed also an
 254 averaged relationship able to estimate the chloride bound by an HCP (Figure 1a). The
 255 proposed function is the sum of the two isotherms described in sections 3.2.1 and 3.2.2
 256 weighted by the corresponding mass fractions

$$257 \quad C_b^{HCP} = K_{CSH} \frac{a_{CSH} C_f}{1 + a_{CSH} C_f} M_{CSH} + K_{AFm} C_f^\alpha M_{AFm}, \quad (21)$$

258 where M_{CSH} and M_{AFm} are the mass fractions of the CSH and AFm .

259 4. Numerical procedure

260 In the present work, the field equation (i.e., the mass balance in Equation 1) is discretized
 261 in space by means of the finite difference method (FDM) using a central scheme in a
 262 3D domain. The geometry is discretized along each direction using a regular mesh of
 263 points with constant step size δ .

264 Since in the present work the behavior of heterogeneous microstructures of HCP
 265 is studied, it is convenient to introduce a quantity that macroscopically can describe
 266 the chloride diffusion behavior of the whole cement matrix at a larger scale than that
 267 of its single phases. This is equivalent to treating the cement paste as a homogeneous
 268 material with a homogenized diffusivity \mathcal{D}_{eff}^H that accounts for the chloride binding.
 269 The homogenized diffusivity should be defined so as to minimize the differences in
 270 behavior between the ideally homogeneous material and the heterogeneous HCP. In
 271 the present work a least square minimization approach is adopted and the transition
 272 from the heterogeneous microscale to the homogeneous material is expressed using the
 273 following objective function

$$274 \quad \Pi := [\langle \mathbf{J} \rangle - \mathbf{J}^H (\langle \nabla C_{tot} \rangle)]^2 \rightarrow \min, \quad (22)$$

where $\langle \cdot \rangle$ is the volume averaging operator defined over the heterogeneous material as

$$\langle \cdot \rangle = \frac{1}{V} \int_V \cdot dV, \quad (23)$$

and $\mathbf{J}^H (\langle \nabla C_{tot} \rangle)$ is the flux in the homogenized material, which is assumed dependent on the average gradient of the total concentration through the homogenized constitutive equation

$$\mathbf{J}^H (\langle \nabla C_{tot} \rangle) = -\mathcal{D}_{eff}^H \langle \nabla C_{tot} \rangle. \quad (24)$$

After substituting Equation 24 into Equation 22, the resulting minimization problem permits one to determine \mathcal{D}_{eff}^H by solving the equation $\partial \Pi / \partial \mathcal{D}_{eff}^H = 0$, leading to

$$\mathcal{D}_{eff}^H = -\frac{\sum_i [\langle J \rangle_i \langle \nabla C_{tot} \rangle_i]}{\sum_i [(\langle \nabla C_{tot} \rangle_i)^2]}, \quad (25)$$

where $i = 1, 2, 3$ are the three orthogonal coordinate axes x, y, z .

5. Numerical simulations and comparisons with experimental results

In this section, the results of the numerical simulations are reported. First, the adopted HCP microstructures are described, then the test setups are illustrated. Finally, numerical and experimental results are compared.

5.1. HCP microstructures

Both X-ray CT scans of a real sample and simulated 3D microstructures of HCP are used here. In particular, the real CT scan is obtained from the Visible Cement Dataset (VCD) available from NIST [27, 28] while the simulated microstructures are generated using CEMHYD3D [25, 26]. The CT images from the VCD are able to distinguish only between hydration products, unhydrated residuals and pores (Figure 2a), while in the simulated HCP all the possible different phases produced during the hydration of the cement clinker are recognized. The cement powder is assumed to be composed of spherical particles whose dimensions and distribution are statistically consistent with high resolution 2D scanning electron microscope (SEM) images of the cement powder [25]. The numerical approach allows one to control each parameter of the sample volume, such as the water-to-cement ratio w/c , and the curing time t_{cur} and conditions (i.e., sealed curing or not, curing temperature). Both simulated and real microstructures used here have the same dimensions, $100 \times 100 \times 100 \mu\text{m}^3$, and resolution of $1 \mu\text{m}^3/\text{voxel}$. Concerning the FDM scheme, one node for each voxel is used.

The adopted HCP microstructure from the VCD database [27, 28], which is shown in Figure 2a, comes from the CT scan of a hydrated cement paste prepared using *Cement and Concrete Reference Laboratory Cement 133* or *CCRL133* [35] with $w/c=0.45$ by

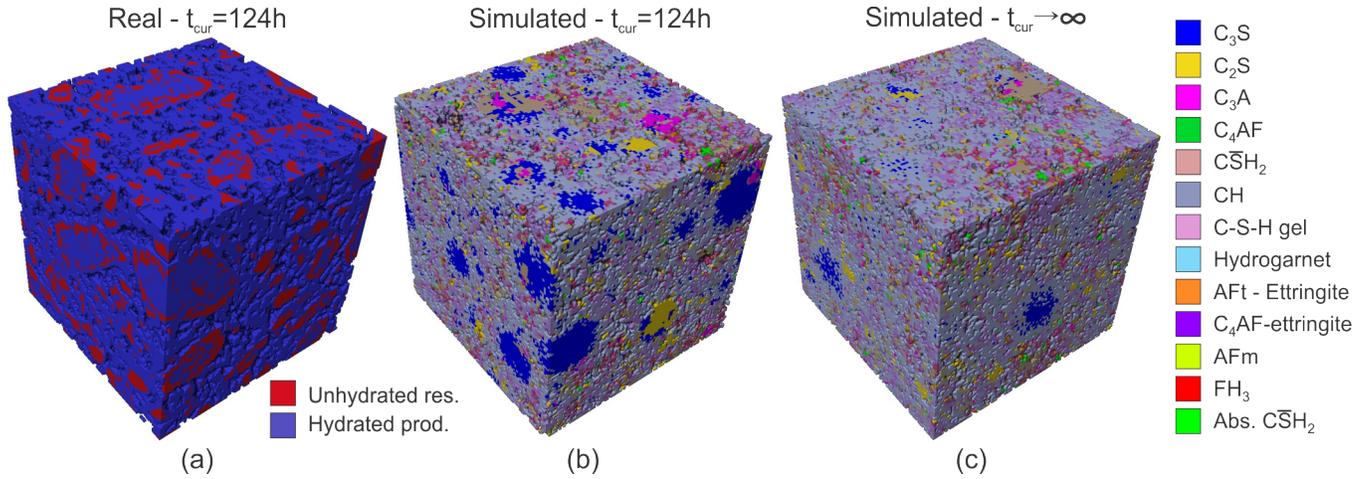


Figure 2. Voxel-based microstructures used in the numerical analyses: (a) real HCP from CT scans available at the VCD website [27, 28], (b) simulated HCP, representing the real HCP from the CEMHYD3D code [25, 26], (c) the same HCP but for an ideally infinite curing time.

307 mass. This is an ordinary Portland cement (OPC) with a standard composition (i.e.,
 308 no additives, fly ashes, limestone or slag particles are present) and is used here for
 309 validation purposes since it is one of the most studied cements in the literature. A
 310 curing time under unsealed conditions of $t_{cur}=124$ h is selected because it is the longest
 311 curing time available in the VCD database [27, 28]. It is worth noting that for a curing
 312 time of 124 h, the hydration degree is about 0.63 and only a limited amount of hydration
 313 products is present (see section 5.2.6), thus the effects of binding on diffusion are not
 314 very pronounced. However, this specimen is used here to confirm that the simulated
 315 HCPs give results similar to real microstructures when studying chloride diffusion and
 316 binding.

317 Concerning the simulated HCPs, two different types of microstructure are used.
 318 One is meant to reproduce the real HCP adopted (i.e., $w/c = 0.45$ and cured under
 319 unsealed conditions for $t_{cur}=124$ h at 20 °C, see Figure 2b). The second type of HCP
 320 simulates a fully cured situation (i.e., $t_{cur} \rightarrow \infty$ in unsealed conditions at a temperature
 321 of 20 °C, see Figure 2c), and is obtained for eight different w/c ratios - namely, 0.35,
 322 0.40, 0.45, 0.50, 0.55, 0.60, 0.70 and 0.80. For usual applications, a w/c ratio in the
 323 range 0.35-0.60 is chosen. Values greater than 0.60 are here adopted only to show the
 324 capabilities of the model. All the initial microstructures are completely flocculated,
 325 i.e. the clinker particles are randomly moved in order to obtain a single cluster of
 326 particles mutually connected (namely, a single floc [26, 36]). This choice is justified
 327 since ordinary cements, such as *CCRL133*, have a strong tendency to flocculate, unless
 328 specific additives are used [36].

329 *5.1.1. Segmentation strategy for the real HCP microstructures* The graylevel threshold
 330 values for the segmentation of the raw CT images are obtained from the graylevel

Table 2. Volume fractions and hydration degrees for the real and simulated HCPs with $t_{cur} = 124$ h.

t_{cur} [h]	HCP type [-]	V_{pores} [Vol. %]	V_{hyd} [Vol. %]	V_{unhyd} [Vol. %]	α_{hyd} [-]	ν_{hyd} [-]
0	Real	59.1	0.0	40.9	0.00	-
0	Simulated	59.3	0.0	40.7	0.00	-
124	Real	28.9	56.3	14.8	0.64	2.16
124	Simulated	30.1	55.3	14.6	0.64	2.12

331 histogram using as a constraint the degree of hydration $\alpha_{hyd}(t)$ and the average cement
 332 stoichiometry. The degree of hydration measures the amount of cement reacted and is
 333 defined as

$$334 \quad \alpha_{hyd}(t) = \frac{V_{clink} - V_{unhyd}(t)}{V_{clink}}, \quad (26)$$

335 where $V_{unhyd}(t)$ is the volume fraction of unhydrated residuals at a certain time t and
 336 V_{clink} is the initial volume of clinker (i.e., $V_{clink} = V_{unhyd}(0)$). To account for the cement
 337 stoichiometry, the following hydration expansion factor $\nu_{hyd}(t)$ is used

$$338 \quad \nu_{hyd}(t) = \frac{V_{hydr}(t)}{V_{clink} - V_{unhyd}(t)}, \quad (27)$$

339 where $V_{hydr}(t)$ is the volume fraction of the hydrated products at the time t .

340 The segmentation thresholds are chosen in such a way as to obtain a ν_{hyd} of about
 341 2.15, which is an average value for an OPC [26, 37], and to reproduce the degree of
 342 hydration $\alpha_{hyd}(t)$ as deduced by the CEMHYD3D simulated microstructure. The two
 343 parameters α_{hyd} and ν_{hyd} for the real and simulated microstructures are reported in
 344 Table 2, along with the volume fractions of pores, unhydrated residuals and hydrated
 345 products.

346 The validity of the segmentation based on the above criterion has been checked
 347 by means of the widely used Power's model [37, 38], which allows one to estimate the
 348 volume fractions of pores V_{pores}^P , unhydrated residuals V_{unhyd}^P and hydrated products V_{hyd}^P
 349 as follows

$$350 \quad V_{pores}^P = \frac{w/c - 0.36\alpha_{hyd}}{w/c + 0.32},$$

$$350 \quad V_{unhyd}^P = \frac{0.32(1 - \alpha_{hyd})}{w/c + 0.32}, \quad (28)$$

$$V_{hyd}^P = 1 - (V_{pores}^P + V_{unhyd}^P).$$

351 As observable comparing Table 2 and 3 the values obtained with both CEMHYD3D
 352 and through the segmentation of the real CT images are in very good agreement with
 353 each other and with the values estimated from Power's model.

Table 3. Volume fractions for the real and simulated HCPs with $t_{cur} = 124$ h as deduced using Power’s model.

t_{cur} [h]	HCP type [-]	V_{pores}^P [Vol. %]	V_{hyd}^P [Vol. %]	V_{unhyd}^P [Vol. %]
124	Real	28.8	56.0	15.2
124	Simulated	28.1	57.3	14.6

Table 4. Volume fractions and hydration degrees of the tested HCPs with $t_{cur} \Rightarrow \infty$.

w/c [-]	V_{pores} [Vol. %]	V_{hydr}		V_{unhydr} [Vol. %]	α_{hyd} [-]
		Binding [Vol. %]	No binding [Vol. %]		
0.35	14.3	45.8	27.8	12.1	0.74
0.40	15.9	48.4	28.5	7.2	0.84
0.45	18.5	49.8	28.0	3.7	0.91
0.50	22.4	47.3	27.4	2.9	0.92
0.55	26.5	44.9	26.1	2.5	0.93
0.60	30.9	42.1	24.3	2.7	0.92
0.70	37.6	38.0	22.4	2.0	0.93
0.80	42.6	35.4	20.4	1.6	0.94

354 *5.1.2. Effect of the w/c ratio on porosity and pores topology* The increment of the
 355 w/c ratio is related to a higher amount of voids, as visible in Table 4, where the volume
 356 fractions for all the HCPs are listed, and in Figure 3, where the distributions of hydrated
 357 products and unhydrated residuals are also reported for different HCPs (namely, for
 358 $w/c = 0.35-0.60-0.80$). Figure 3 also shows that, passing from $w/c = 0.35$ to 0.60, the
 359 voids network is more extended but topologically similar, meaning that it keeps the
 360 characteristic of capillarity (i.e., the presence of small voids connected by capillaries).
 361 Conversely, for the HCP with $w/c = 0.80$ an alveolar-like structure characterized by
 362 large cavities forming preferential pathways for the diffusion is present. However, these
 363 cavities are connected by capillaries that obstruct ions migration.

364 *5.1.3. Percolation of the pore network* Another factor strongly related to the diffusive
 365 behavior is the percolation of the pore network (see [7, 8] for a detailed review). A
 366 system of voids is termed percolated in a certain direction if, in that direction, there
 367 exists at least one continuous path that goes from one end to the opposite one. If such
 368 a path does not exist, the pore network is said to be depercolated. In the latter case,
 369 the ions should overcome membranes of solid material that isolate the capillary pores
 370 in clusters, delaying the diffusion [8]. Garboczi and Bentz [8] demonstrated that the
 371 depercolation limit for a Portland HCP corresponds to a void volume fraction of 18 %,

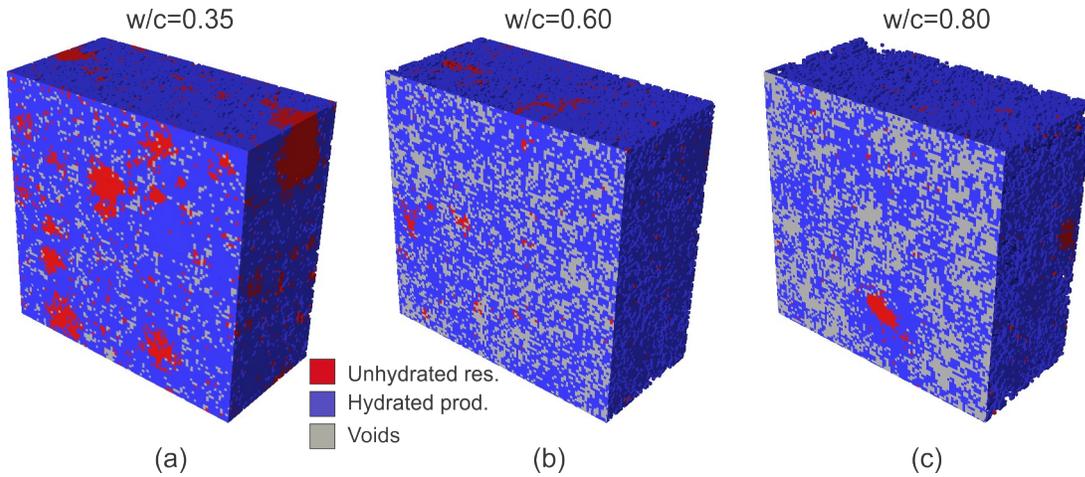


Figure 3. Sections of the simulated microstructures used for the diffusion test showing distribution of hydration products, unhydrated residuals and voids: (a) $w/c=0.35$, (b) $w/c=0.60$, (c) $w/c=0.80$.

372 using a pore resolution of $1 \mu\text{m}$ in the model. Above this limit the probability to find
 373 a percolated pore system in an ideally infinite domain is 100 %, while below this limit
 374 such probability is 0 %. In this latter case, the HCP microstructure contains many
 375 depercolated subpaths and isolated clusters of pores. The former are pore systems that
 376 can be directly accessed from one of the outer exposed surfaces, while the isolated pores
 377 can be reached by diffusing species only after passing through a region of solids. The
 378 extent in volume of the percolated, depercolated and isolated pores, the presence of
 379 bottlenecks in the percolation path and its tortuosity strongly influence the diffusive
 380 behavior of the HCP [10].

381 The changes in the capillary pore network are observable in Figure 4, where the
 382 evolution of the percolated and depercolated pore systems for the HCPs with $w/c=0.40$ -
 383 0.45 - 0.50 is examined using a burning algorithm similar to the one proposed by Bentz
 384 and Garboczi in [7]. In particular, it is shown that, for the HCPs used here, 0.45 is
 385 the lowest w/c that produces a percolated path, which however is limited in extension
 386 and characterized by the presence of bottlenecks and a high tortuosity (Figure 4b).
 387 Conversely, the microstructure with $w/c=0.40$ is depercolated (Figure 4a) and the one
 388 with $w/c=0.50$ has already a very large percolated system.

389 Figure 5 reports the percolated, depercolated and isolated fractions of the total
 390 pore volume as functions of the w/c ratio for fully cured systems. It can be seen how
 391 the pore connectivity changes with the w/c ratio, confirming that below a pore volume
 392 of nearly 18 % (i.e., for $w/c \leq 0.40$) the microstructures are depercolated while for
 393 $0.45 \leq w/c \leq 0.60$ the extension of the percolated system grows rapidly, then for $w/c >$
 394 0.60 the curve flattens until reaching a value of about 93 % at $w/c=0.80$. The afore-
 395 mentioned burning algorithm is applied also to the samples with $t_{cur}=124$ h resulting
 396 in a percolated void fraction of 79.8 % and 99.1 % respectively for the simulated and
 397 real microstructure. These results are obviously higher than their fully hydrated coun-

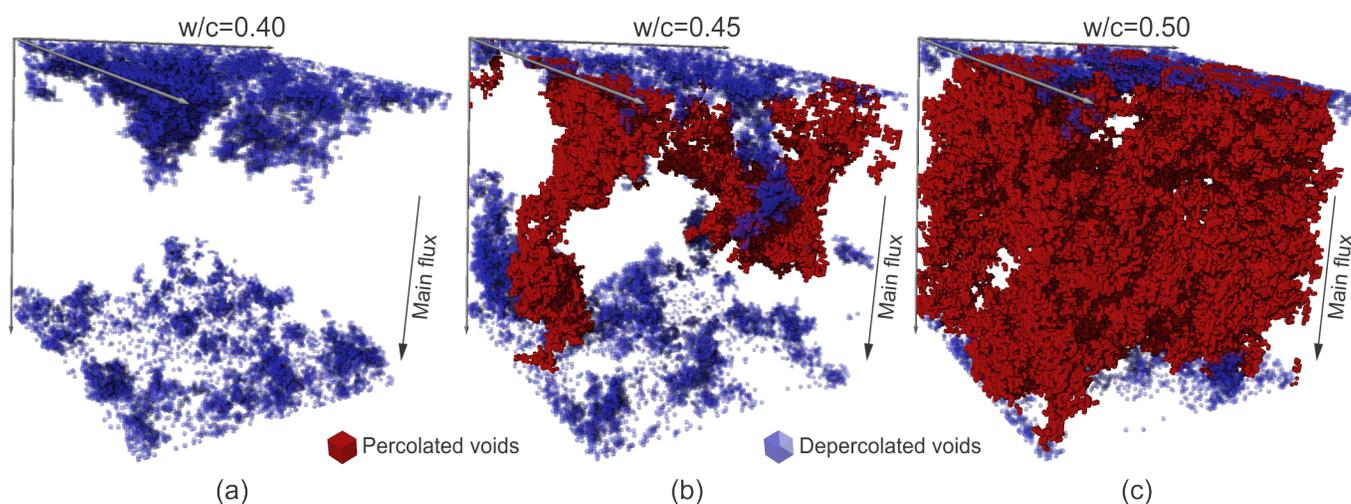


Figure 4. Percolated and depercolated pathways determined using a burning algorithm for different HCPs: (a) $w/c=0.40$, (b) $w/c=0.45$ and (c) $w/c=0.50$.

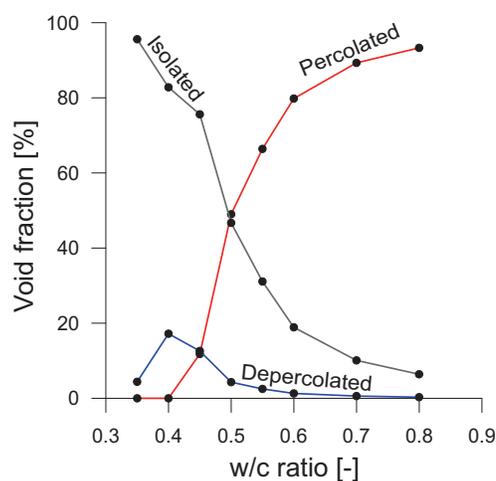


Figure 5. Percolated, depercolated and isolated fractions of the total pore volume.

terparts because of the limited curing time, however they are in good agreement with those reported in [37] for a similar case. Although it is recognized that the chosen voxel resolution will influence both the percolation of the capillary porosity and the resulting diffusivity [39], in this study, the resolution for the simulated microstructures was set to $1 \mu\text{m}^3/\text{voxel}$, following the results of previous studies where simulated microstructures are compared to experimental CT images [40]. Furthermore, the adopted resolution matches the one of real microstructures of the VCD database [27, 28].

5.1.4. Comparison between real and simulated HCPs In Figure 6 the real and simulated microstructures of the non-hydrated cement clinker used here are compared and the corresponding volume fractions are reported in Table 2. Although the real geometry is composed of highly irregular clinker particles while the simulated ones are spherical,

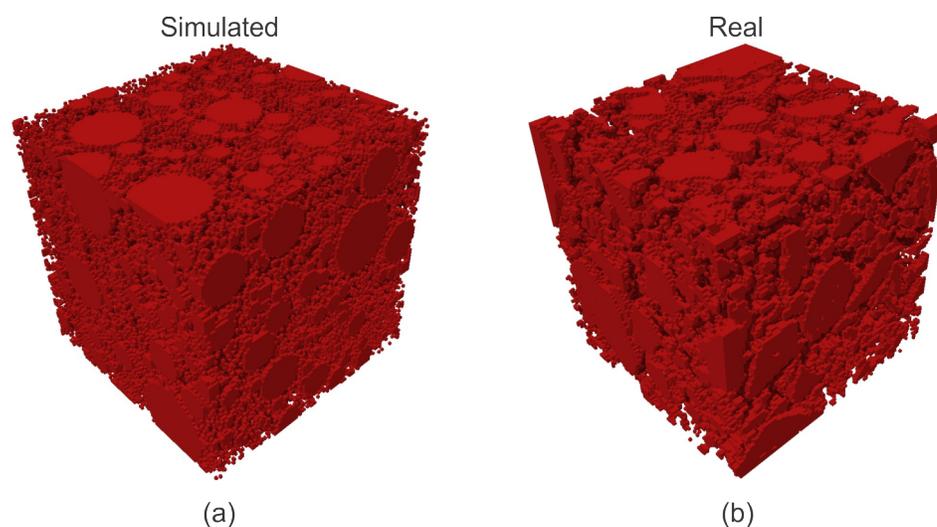


Figure 6. Clinker (non-hydrated cement powder) 3D microstructures: (a) simulated using CEMHYD3D [25, 26], (b) from CT scan images (from VCD [27, 28]).

409 a good agreement is noticeable, especially concerning the presence and distribution of
 410 both large and small clinker particles.

411 Similar conclusions can be drawn comparing the real and simulated (hydrated)
 412 HCPs depicted in Figure 7. In particular, as also reported in [37], the microstructures
 413 appear similar even though the geometry coming from the CT scan images (Figure 7b)
 414 is more irregular than the simulated counterpart (Figure 7a). However, for the real
 415 HCP only a coarse subdivision of the phases is available (i.e., voids, hydrated products
 416 and unhydrated residuals) because of the limits of the 3D imaging techniques available
 417 at that date. Conversely, although in the simulated microstructures the geometry is
 418 idealized, all the different phases present in HCP are finely resolved with a precision of
 419 $1 \mu\text{m}^3$. Moreover, the simulation model used [25, 26] allows one to directly and easily
 420 control all the principal parameters of a cement clinker and of a HCP.

421 In Table 2 the volume fractions of voids, hydrated products and unhydrated
 422 residuals for the real and simulated microstructure used herein are reported. The
 423 marginal differences in the values are possibly related to the fact that the w/c ratio
 424 in the simulated microstructures is directly enforced on the tested sample, while in the
 425 real HCP it is imposed on the whole volume from which the sample is extracted, that
 426 is much larger than $100 \times 100 \times 100 \mu\text{m}^3$.

427 In [37], for a cement similar to the one analyzed here, hydrated and clinker
 428 microstructures from CT scan images and generated via CEMHYD3D adopting
 429 both spheres and real irregular particles are compared. The author demonstrated
 430 that the employment of spherical geometries does not affect the phase distribution
 431 in microstructures with the same w/c ratio, provided that a reliable particle size
 432 distribution is given [37].

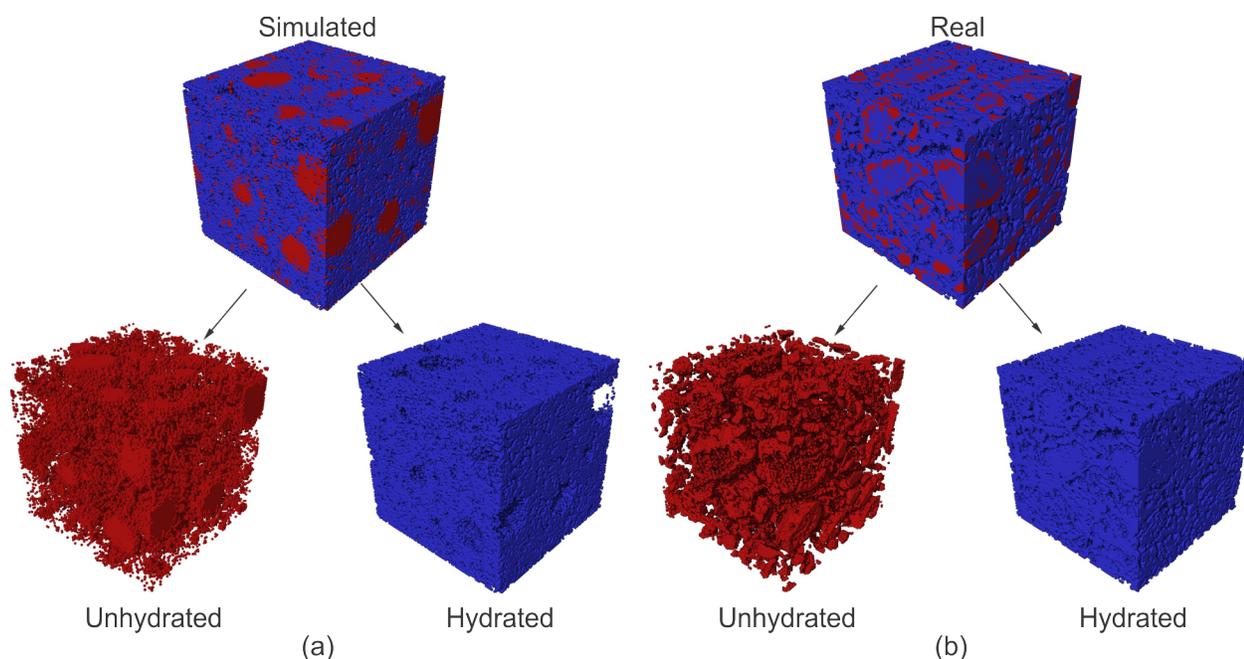


Figure 7. HCPs with $t_{cur}=124$ h used in the numerical simulations and their hydrated products and unhydrated residuals: (a) simulated using CEMHYD3D [25, 26], (b) from CT scan images (from VCD [27, 28]).

433 5.2. Numerical simulations of diffusion and binding

434 The diffusion test is performed by prescribing a concentration gradient between two
 435 opposite faces of a sample. By tracking the amount of chloride ions reaching the outlet
 436 face, the diffusion coefficient of the HCP is estimated [29, 41, 42]. The most classic
 437 setup involves an HCP sample where at one face (inlet face) a chloride concentration
 438 C_{inlet} is applied, while the opposite face (outlet face) is in contact with a solution with
 439 concentration C_{outlet} equal to zero (i.e., pure water). All the other faces are sealed, i.e.
 440 the chloride flux normal to the surface is zero. Differently, the binding test involves the
 441 application of a fixed time-invariant concentration of chlorides on all the faces of the
 442 specimen (i.e., the sample is immersed in a chloride bath) and is used to test the binding
 443 capability of the HCP [11]. By testing several samples at different concentrations, the
 444 isotherm curve for the tested HCP is obtained.

445 In the following, after the definition of the parameters adopted in the performed
 446 analyses, the numerical results are compared to experimental results for the diffusion
 447 tests of Page et al. [29], Yu and Page [41], Nagala and Page [43], MacDonald and
 448 Northwood [44] and Tang and Nilsson [42] along with the binding tests of Hirao et al.
 449 [11]. Also, some observations arising from the comparison between real and simulated
 450 HCPs are drawn.

451 5.2.1. Diffusion coefficients of the HCP phases As already outlined in section 2.1, the
 452 proposed model accounts for the heterogeneity of the values of the diffusivity in the

453 HCP microstructure. Therefore, the diffusivity parameter in Equation 2 (and, hence, in
454 Equation 4) is here written as

$$455 \quad D(\mathbf{x}) = \sum_h D_h \delta_{h,k(\mathbf{x})}, \quad (29)$$

456 where the summation is evaluated over all the possible phases h with diffusivity D_h
457 present in the microstructure, $k(\mathbf{x})$ is the phase at the point \mathbf{x} , and $\delta_{h,k(\mathbf{x})}$ is the
458 Kronecker delta evaluated at that point, which is equal to 1 if $k(\mathbf{x}) = h$ and 0 otherwise.
459 Because of the reduced dimension of the voids in the HCP (transparent in Figure 2), a
460 value for the diffusivity of chloride ions in capillary pores (see Section 2) $D_c = D_{pores} =$
461 $3.0 \cdot 10^{-10} \text{ m}^2/\text{s}$ at room temperature ($T=20^\circ\text{C}$) is assumed, in very good agreement
462 with values proposed by other authors [17, 29]. Conversely, for all other phases the
463 solid diffusivity D_s should be adopted. In this case the diffusivity is much lower than
464 in pores because, although in the solid material a capillary network is still present
465 because of the agglomeration of crystals, the diameter of the micropores is much smaller
466 and the tortuosity is higher. Moreover, D_s accounts also for the part of the volume
467 effectively occupied by the crystal grains. For the diffusivity of the phases exhibiting
468 binding (*CSH* and *AFm*), the well accepted empirical formula proposed by Garboczi
469 and Bentz [8] $D_{bind} = D_{AFm} = D_{CSH} = D_{pores}/400$ is adopted here for both *CSH* and
470 *AFm*, although it is recognized that they are two different hydration products. This
471 assumption is needed since it is not possible to find in the literature a reliable exper-
472 imental estimation of such parameters. According to the literature, all other hydrated
473 or unhydrated phases show a lower diffusivity than these binding phases, hence here
474 a value of $D_{NObind} = D_{unhydr} = D_{bind}/10 = D_{pores}/4000$ is assumed in order to ac-
475 count for the presence of nanopores smaller than a fraction of micron. The nanopore
476 network arises from the imperfect coalescence of different grains of the solid phase.
477 Concerning the real microstructure, D_{unhydr} is associated with all unhydrated residuals,
478 while for the hydration products it is assumed that their volumetric composition is
479 voxel-wise equal to the hydration products composition of the simulated HCP, which
480 is composed at $t_{cur}=124$ h by 59.5 % of *CSH* and by 1.0 % of *AFm* while the
481 remaining part is composed of non-reactive phases. Then, following the additivity of
482 Equation 29, the hydrated diffusivity is taken as the weighted average of the respective
483 diffusivities, i.e. $D_{hydr} = (0.595 + 0.010)D_{bind} + (1 - 0.605)D_{NObind}$. The same approach
484 is used to define the binding curve of the hydration products in the real HCP, namely
485 $C_b^{hydr} = 0.595C_b^{CSH} + 0.010C_b^{AFm}$, while the binding coefficient follows the relationship
486 of Equation 12. All the adopted parameters are summarized in Table 5. The low amount
487 of *AFm* crystals is justified by the short curing time (124 h). In CEMHYD3D the *AFm*
488 phase forms from the ettringite products (*AFt*) after the complete sulphate depletion
489 [25, 26], hence the *AFm* volume fraction increases by about a factor of 3 for a fully
490 hydrated microstructure (Sect. 5.2.5).

491 5.2.2. Numerical simulation of the diffusion tests Two series of tests are here
492 reproduced whose main difference is the imposed chloride concentration at the inlet

Table 5. Diffusion coefficients used in the numerical simulations.

D_{pores} [$10^{-12} \text{ m}^2/\text{s}$]	$D_{bind} = D_{CSH} = D_{AFm}$ [$10^{-12} \text{ m}^2/\text{s}$]	D_{NObind} [$10^{-12} \text{ m}^2/\text{s}$]	D_{unhydr} [$10^{-12} \text{ m}^2/\text{s}$]	D_{hydr} [$10^{-12} \text{ m}^2/\text{s}$]
300.000	0.750	0.075	0.075	0.480

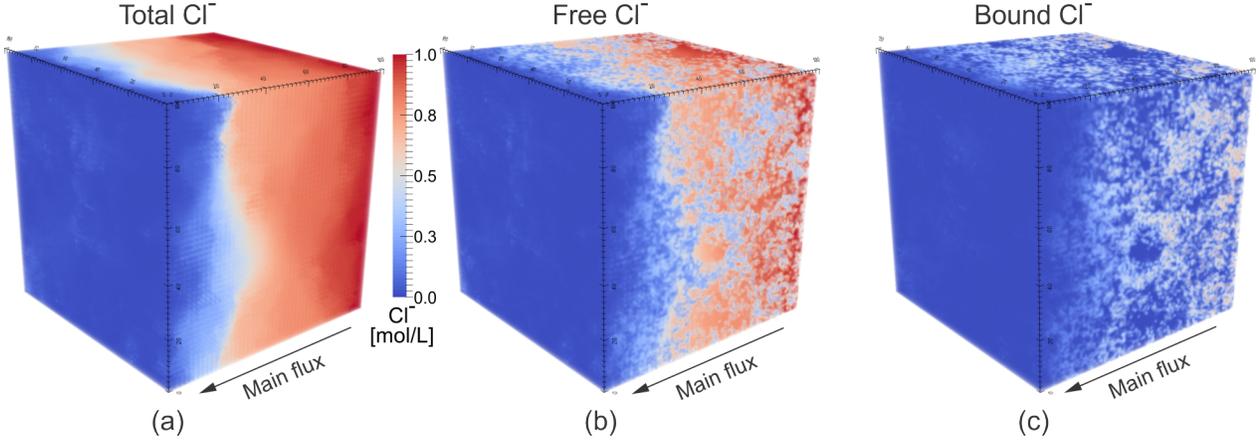


Figure 8. Steady-state chloride concentrations from the numerical simulation of the Page et al. [29] test for the simulated HCP with $t_{cur} \rightarrow \infty$ and $w/c=0.45$: (a) total chloride concentration, (b) free chloride concentration, (c) bound chloride concentration.

493 face, which is $C_{inlet}=1.0 \text{ mol/L}$ for Refs. [29, 41, 43, 44] and $C_{inlet}=0.5 \text{ mol/L}$ for Ref.
 494 [42]. For both, the outlet concentration is $C_{outlet}=0 \text{ mol/L}$ and the flux normal to all
 495 the other faces is set to zero.

496 In Figure 8, the results for $C_{inlet}=1.0 \text{ mol/L}$ in terms of total, free and bound
 497 chloride concentrations for the specimen with $w/c=0.45$ and $t_{cur} \rightarrow \infty$ are presented,
 498 while the results for the other specimens are qualitatively very similar. It is evident that,
 499 although the total chloride concentration distribution appears quite smooth (Figure 8a),
 500 the opposite is true for the free and bound chloride distributions (Figure 8b-c). This is
 501 due to the heterogeneous distribution of the hydration products (Figures 2c) and to the
 502 presence in the clinker structure of large particles that are hydrated only in their outer
 503 part, leading to the formation of clusters of unhydrated residuals as visible in the side
 504 face of the HCP in Figure 8b-c.

505 The results of all the diffusion test simulations in terms of homogenized diffusivity
 506 both accounting for and neglecting the binding effects, respectively \mathcal{D}_{eff}^H and \mathcal{D}_{NObind}^H ,
 507 are reported in Table 6 and illustrated in Figure 9. Moreover, in Figure 10 the influence
 508 of the binding effect on the homogenized diffusivity \mathcal{D}^H is expressed through the ratio
 509 ρ_D defined as

$$510 \quad \rho_D = 1 - \frac{\mathcal{D}_{eff}^H}{\mathcal{D}_{NObind}^H}. \quad (30)$$

Table 6. Results of the numerical simulations of the diffusion tests.

HCP type	w/c	t_{cur}	\mathcal{D}_{eff}^H		\mathcal{D}_{NObind}^H
	[–]	[h]	[$10^{-12} m^2/s$]	[$10^{-12} m^2/s$]	[$10^{-12} m^2/s$]
			0.5 mol/L ^(a)	1.0 mol/L ^(a)	
Simulated	0.35	$\rightarrow \infty$	0.46	0.52	1.12
Simulated	0.40	$\rightarrow \infty$	0.70	0.79	1.46
Simulated	0.45	$\rightarrow \infty$	1.17	1.30	2.29
Simulated	0.50	$\rightarrow \infty$	2.97	3.14	4.50
Simulated	0.55	$\rightarrow \infty$	6.94	7.15	8.62
Simulated	0.60	$\rightarrow \infty$	13.42	13.67	15.22
Simulated	0.70	$\rightarrow \infty$	28.35	28.54	30.07
Simulated	0.80	$\rightarrow \infty$	44.26	44.42	45.66
Real CT scan	0.45	124	-	17.90	18.70
Simulated	0.45	124	-	13.09	14.31

^(a) Values related to the inlet concentration C_{inlet} .

511 From the numerical results in Figure 9 it is possible to notice how an increase
 512 of the w/c ratio leads to an exponential increase of the homogenized diffusivity that
 513 approaches a linear trend for $w/c > 0.6$ (Figure 9 and Table 6). For very high w/c
 514 ratios the homogenized diffusivity \mathcal{D}^H grows rapidly approaching D_{pores} , which is the
 515 expected value in the ideal case of $w/c = \infty$. As already reported by Garboczi and
 516 Bentz [8], such non-linear trend is related to the increment, with the w/c ratio, of the
 517 volume of the capillary pores (see Table 4 and section 5.1.2), which have a diffusivity
 518 value 2-3 orders of magnitude higher than that of the other phases (Table 5). Hence,
 519 even a relatively small increase in pore volume can result in a relevant increment of the
 520 homogenized diffusivity.

521 As already outlined in section 5.1.2, the increment of porosity leads to a change in
 522 the pores topology that, interestingly, takes place in correspondence of the beginning of
 523 the linear branch of the $\mathcal{D}^H - w/c$ curve (Figure 9), i.e. for $w/c \simeq 0.60$ for the cement
 524 used here.

525 The sudden change in slope of the $\rho_D - w/c$ curve for $w/c \simeq 0.45$ (Figure 10)
 526 can be explained considering the percolation properties of the HCP microstructures
 527 (section 5.1.3). Table 4 and Figure 5 show that such slope change takes place
 528 in correspondence of the transition from the percolated to the depercolated HCPs
 529 (Figure 4). Depercolation leads to a dramatic reduction of the pore network connectivity
 530 delaying the penetration of chloride ions. Moreover, as visible in Figure 3a and in
 531 Table 4, for low w/c ratios a larger amount of unhydrated residuals are still present
 532 especially in correspondence of large clinker particles. In such cases, part of the cement
 533 does not react with water leading to the formation of inclusions of unhydrated products,
 534 which are assumed not to bind chlorides.

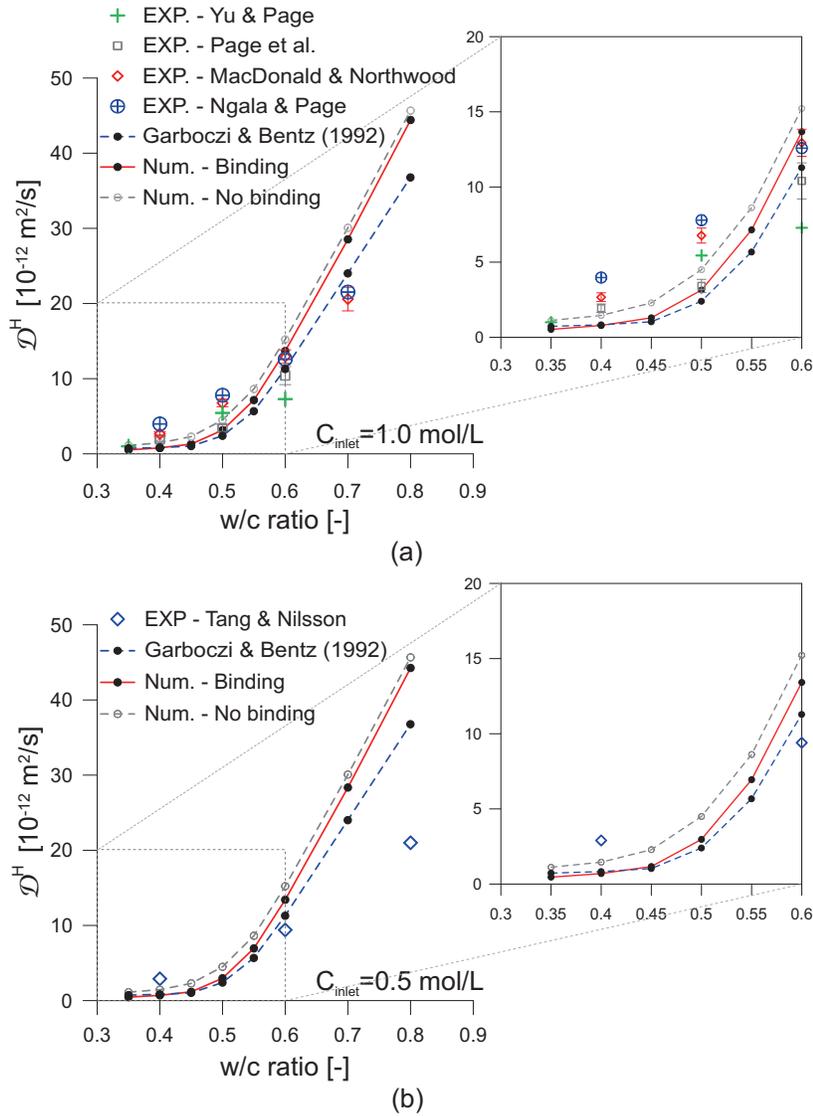


Figure 9. Homogenized diffusivity vs. w/c ratio curves from the numerical analyses of the diffusion tests: (a) tests with $C_{inlet} = 1.0 \text{ mol/L}$, (b) tests with $C_{inlet} = 0.5 \text{ mol/L}$.

535 Although in Figure 9 the difference between homogenized diffusivity with and
 536 without binding is apparently limited, the parameter ρ_D plotted in Figure 10 shows
 537 that the interaction between solid phases and chloride ions indeed plays a significant
 538 role in the diffusive behavior. For the values of w/c commonly employed (i.e. in the
 539 range 0.35-0.60), a reduction of the homogenized diffusivity in the range 10 %-60 % is
 540 observed. Such a relevant influence is due to the small values assumed by the binding
 541 activity coefficients μ_B within the HCP volume in the phases exhibiting binding, as also
 542 illustrated in Figure 11.

543 Figure 10 and Table 6 show that for $w/c > 0.60$ the difference between homogenized
 544 diffusivity with and without binding is less than 10 % even though the solid phase is
 545 mainly composed of hydration products exhibiting binding, as also reported in Table 4.

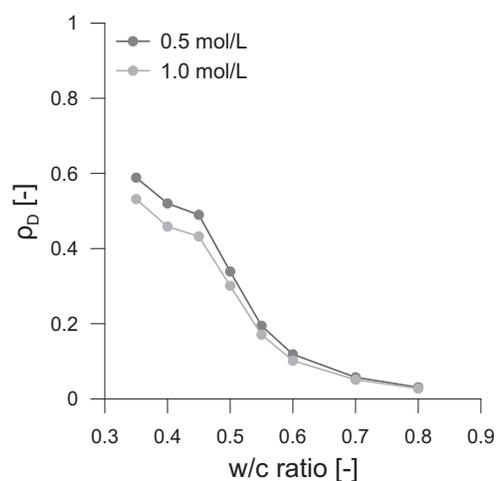


Figure 10. Influence of the binding effect on the homogenized diffusivity (ρ_D ratio) for varying w/c ratio from the numerical analyses of the diffusion tests.

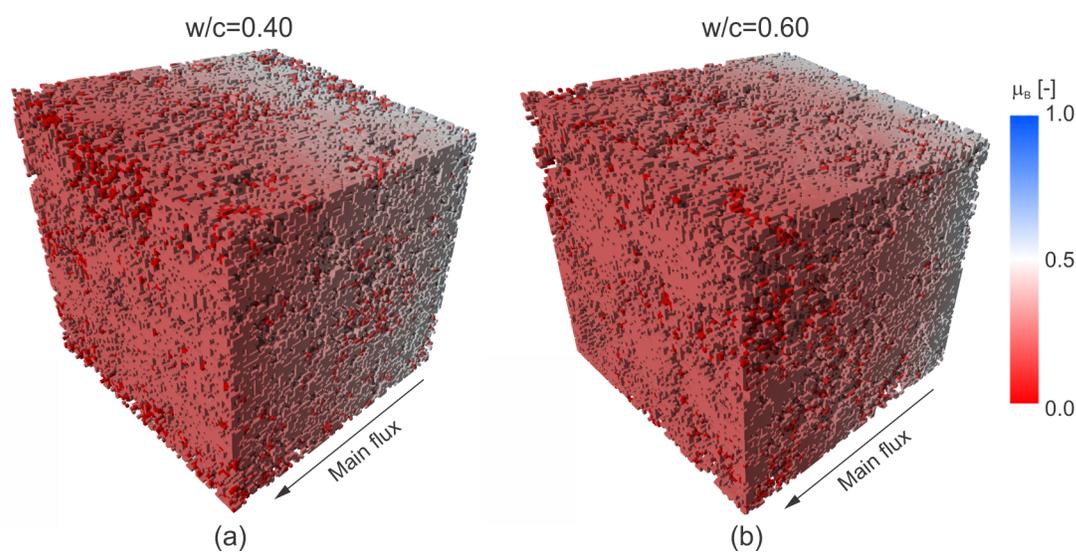


Figure 11. Maps of the calculated diffusivity activity coefficient μ_B for the phases showing binding for the simulated microstructures with (a) $w/c=0.40$ and (b) $w/c=0.60$.

546 This is due to the decreased volume of solid phase with respect to the capillary pore
 547 network, whose volume fraction is well above the percolation limit [8]. In such cases the
 548 alveolar-like void structure dominates the diffusion process because of the high pores
 549 diffusivity and, at the same time, the global effect of binding is reduced since binding
 550 does not take place in water saturated pores. This leads to a drastic change in the
 551 diffusion process, confirming the preminent role of the voids in the diffusion process for
 552 HCPs with a volume fraction of capillary pores much higher than the percolation limit
 553 [8].

554 The influence of binding is less pronounced in the specimens with limited curing

time $t_{cur}=124$ h than for fully hydrated pastes (see Table 6) because the progression of the hydration process implies more hydration products exhibiting binding and, thus, its effect becomes more prominent in later age specimens, as will be also illustrated in section 5.2.6. Moreover, the value of the homogenized diffusivity appears inversely related to the curing time (Table 6). This happens because the hydration process reduces drastically the amount of capillary voids in HCP and, as already noted, increases the amount of hydration products that exhibit binding. These two effects both lead to a reduction of the diffusivity with further hydration.

5.2.3. Comparison with the experimental results of the diffusion tests Figure 9 also compares the numerical results with the experimental data [29, 41–44]. A reasonable agreement is observable between the test results and the numerical curves accounting for binding for both $C_{inlet}=1.0$ mol/L (Figure 9a) and $C_{inlet}=0.5$ mol/L (Figure 9b). However, the difference between the cases with and without binding is very close to the deviation between the various sets of the experimental results. The latter deviation appears quite high, but this is not too surprising considering that the various series of tests used involve different types of cement clinker, temperatures, specimen dimensions and test setups. Such effects may significantly modify the value of the pore diffusivity D_{pores} and, hence, all the diffusivity parameters. Unfortunately, the literature does not report, to date, enough information to allow for a precise and detailed calibration of the diffusivity parameters accounting for all the possible phenomena influencing diffusion. Hence, as also mentioned in section 5.2.1, the parameters adopted here are not finely tuned for the cases at hand. Furthermore, the values of D_{pores} present in the literature are usually deduced neglecting the binding effect, most of the time trying to best fit a simplified solution of Fick’s equation with a set of experimental results. It is finally worth noting that, accounting only for the range of w/c ratio commonly used (i.e., $w/c < 0.60$) and for data from the same experimental campaign, the deviation becomes much smaller. **Certainly a proper case-wise calibration of the parameters would give better agreement with a specific dataset but this is not the goal of the present work. The present comparison should be conceived as a general validation aimed at demonstrating that the model proposed is able to capture, adopting both simulated and real microstructures, the main processes governing diffusion at the microscale as well as the effects of the main variables involved.**

As expected, the numerical results in the range $0.35 \leq w/c < 0.60$ always lie below the experimental points. This behavior is reasonable considering that the experimental tests were performed with $t_{cur} < \infty$ and that the homogenized diffusivity \mathcal{D}^H decreases with the progress of the hydration. From Figure 9 it is also evident that, for very porous HCPs (i.e., for $w/c \geq 0.60$), the homogenized diffusivity is overestimated. However, heterogeneous specimens are likely to be produced for so highly porous HCPs, because of the presence of large cavities inducing phenomena such as bleeding and/or sedimentation. Also, as noted before, for microstructures with $w/c > 0.60$ the diffusion mechanism can be completely different from Fickian diffusion because of the changes in

596 the topology and percolation characteristics of the capillary pore network (Figures 3, 4
597 and 5). In particular, the increment in pores dimension might promote the presence of
598 convective contributions to the particles motion.

599 **It is worth noting that, following what reported in [11], there is the possibility**
600 **that, under particular conditions, also the *Aft* (ettringite) reacts with chloride ions**
601 **producing Friedel's salt following a mechanism similar to AFm. Accounting for the *Aft***
602 **products as reactive phases with a binding isotherm given as in Eq. 19 the changes in**
603 **the final results are limited, in the order of about 0.5-1 %. However, the role of *Aft* in**
604 **the binding process is still controversial and is not accounted for henceforth.**

605 The aforementioned different behavior for porous HCPs is also implicitly considered
606 in the semi-empirical relationship based on the percolation theory proposed by Garboczi
607 and Bentz [8] that, according to the authors, can be adopted to characterize HCPs with
608 $w/c \leq 0.60$. The formula, which is reported for comparison in Figure 9, reads

$$609 \frac{\mathcal{D}^H}{D_{pores}} = 10^{-3} + 0.07 \left(\frac{V_{pores}}{100} \right)^2 + h(\xi) [1.8\xi^2] \quad \text{with} \quad \xi = \frac{V_{pores}}{100} - \frac{V_c}{100}, \quad (31)$$

610 where all the volume fractions are expressed in percentage, $V_c = 18\%$ is the percolation
611 threshold for an OPC [8] and $h(\xi)$ is the Heaviside step function, which is equal to 0
612 when $\xi \leq 0$ and 1 if $\xi > 0$.

613 The obtained numerical results are generally in good agreement with Equation 31
614 (Figure 9) confirming the reliability of the model proposed and the leading role of
615 the pore percolation. In particular, note that the range of validity of Equation 31
616 corresponds to the range of w/c ratios where the proposed approach is more accurate,
617 and, within this applicability range (i.e., for $w/c \leq 0.60$), the agreement between
618 the proposed approach and the experimental data is similar to the one achieved with
619 Equation 31 (Figure 9). However, Equation 31 is based on best fitting procedures
620 and does not directly account for binding and for the heterogeneous distribution of the
621 various phases, meaning that it cannot be generalized.

622 *5.2.4. Comparison with analytical bounds* Figure 12 illustrates the comparison of the
623 numerical and experimental results with analytical homogenization bounds obtained
624 neglecting the binding behavior, including the Voigt, Reuss, Voigt-Reuss-Hill (VRH),
625 Hashin-Shtrikman upper, lower and average bounds (respectively HS+, HS-, Avg. HS).
626 Analytical bounds are very large and also their average values (VRH and Avg. HS curves
627 Figure 12) are usually significantly higher than the experimental results, demonstrating
628 that the adoption of a numerical approach is mandatory to achieve reasonable accuracy.

629 *5.2.5. Numerical simulation of the binding tests* Simulating the binding tests in [11]
630 **(i.e., imposing to all the six faces of the specimen the same concentration)**, it is possible
631 to obtain the numerical binding isotherms, which are reported in Figure 13 for the HCPs
632 with w/c of 0.40 and 0.60. The same plot also contains the average curve proposed
633 by Hirao et al. [11] (Equation 21), where the mass fractions for the two specimens

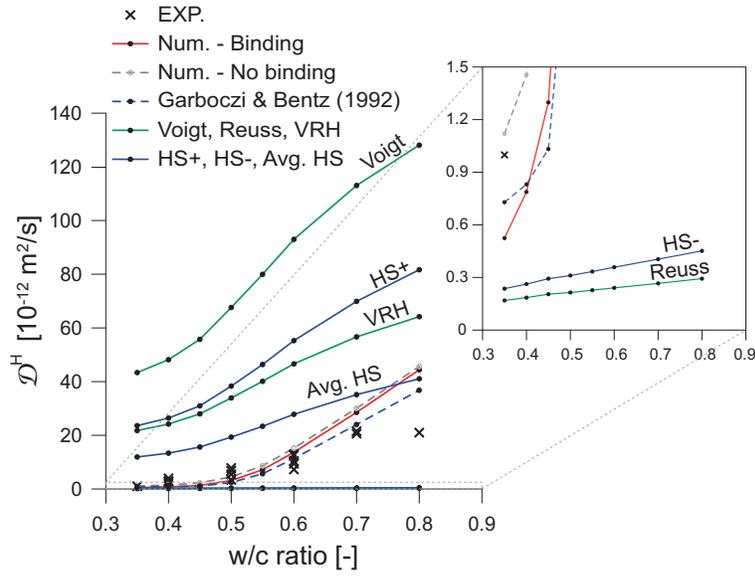


Figure 12. Comparison between the homogenized diffusivity from experimental results, numerical results and various analytical bounds, such as Voigt bound, Reuss bound, Voigt-Reuss-Hill bound (VRH), Hashin-Shtrikman upper, lower and average bounds (respectively HS+, HS- and Avg. HS).

Table 7. Fractions of the binding phases for $w/c=0.40$ and 0.60 .

w/c	AFm		CSH	
	Vol. %	Mass %	Vol. %	Mass %
0.40	2.90	3.25	45.42	53.49
0.60	2.05	2.27	40.11	47.25

634 are estimated from the respective fully cured simulated microstructure (Table 7) using
 635 the phases and HCP characteristics reported in Table 1. The numerical simulations
 636 show a lower amount of bound chloride ions when compared with the prediction of the
 637 empirical averaged relationship of Hirao et al. [11]. These differences can be explained
 638 considering that in Hirao et al. [11] it is implicitly assumed that the concentration of
 639 the free chlorides is uniform in all the phases (voids, hydrated products and unhydrated
 640 residuals) while the numerical curves reported in Figure 13 are the volume average of
 641 the whole HCP, where the free chloride concentration is not homogeneous because of
 642 the effects of the binding. Accounting for the aforementioned issues, the agreement is
 643 acceptable.

644 *5.2.6. Comparison between the diffusive behavior of real and simulated HCP* In
 645 Figures 14 and 15 the contour maps of the total, free and bound chloride concentration
 646 respectively for the simulated and real microstructures with $t_{cur}=124$ h and $w/c=0.45$
 647 are reported. Despite the different starting particle geometry, the two images show a
 648 good agreement, both qualitatively and quantitatively. The slight overestimation of

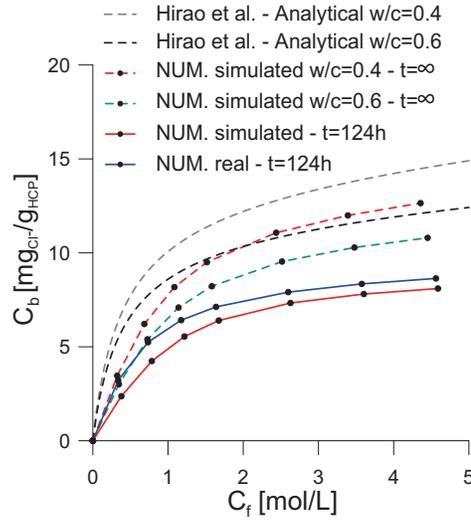


Figure 13. Binding curves from the numerical simulations of the binding tests.

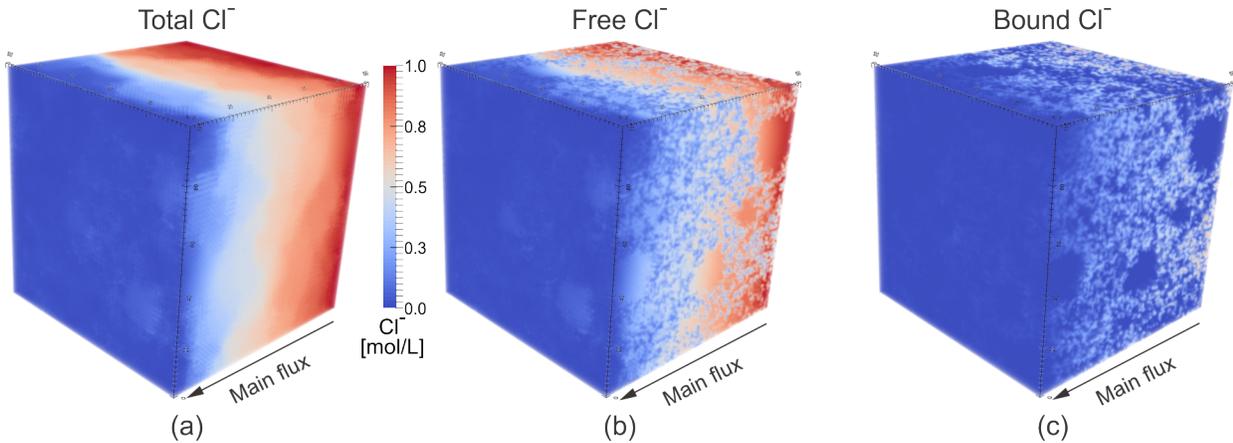


Figure 14. Steady-state chloride concentrations from numerical simulation of the test with $C_{inlet}=1$ mol/L for the simulated HCP with $t_{cur}=124$ h and $w/c=0.45$ reproducing the CCRL133 cement [27, 28]: (a) total chloride concentration, (b) free chloride concentration, (c) bound chloride concentration.

649 the homogenized diffusivity noticeable in Table 6 for the real microstructure agrees
 650 with the differences in the volume fraction of the percolated pore network. However,
 651 such difference is also partly due to the assumed relationship for the diffusivity and
 652 the binding isotherm described in section 5.2.1. In particular, the adopted hydration
 653 products diffusivity is weighted on the volume fractions of the various hydration products
 654 constituting thus an upper bound value (namely, a Voigt bound). Moreover, the binding
 655 isotherm for hydration products is calculated on a volume average basis as well, limiting
 656 thus the possibility to reach values of μ_B close to zero because of the presence of phases
 657 that are insensitive to the binding process.

658 Concerning the binding test Figure 13 demonstrates that the behavior of real and
 659 simulated microstructures is very similar. The difference in the amount of chlorides

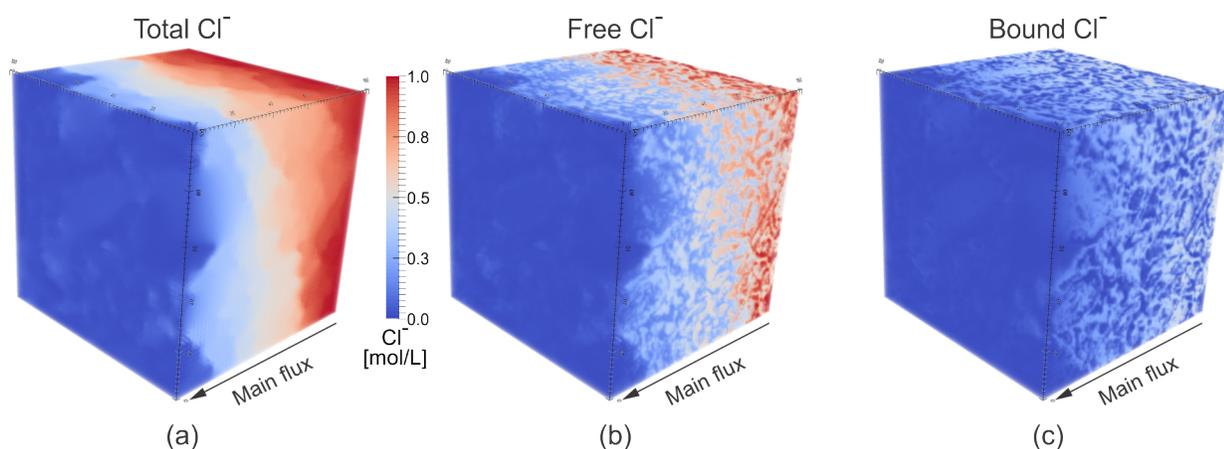


Figure 15. Steady-state chloride concentrations from numerical simulation of the test with $C_{inlet}=1$ mol/L for the real CCRL133 [27, 28] HCP with $t_{cur}=124$ h and $w/c=0.45$: (a) total chloride concentration, (b) free chloride concentration, (c) bound chloride concentration.

660 bound is probably due to the distribution of the phases exhibiting binding, that are
 661 lumped in all the hydrated products in the real microstructure, while they are pointwise
 662 identified in the simulated one (Figure 2).

663 The above evidence makes it possible to conclude that, for the steady-state chloride
 664 diffusion behavior, the results of simulated and real HCP microstructures show a good
 665 agreement provided that for the latter a sufficiently precise estimate of the volume
 666 fractions of the phases exhibiting binding is available.

667 6. Conclusions

668 The problem of diffusion of chloride ions into hardened cement paste (HCP) at the
 669 microscale level is addressed. The microstructures of the HCP are obtained through
 670 computer tomography (CT) images of a real sample as well as simulated using
 671 CEMHYD3D. A model is proposed based on steady-state Fickian diffusion accounting
 672 explicitly for the binding effects through the introduction of a binding activity coefficient
 673 deduced by means of binding isotherms. The model, implemented with the finite
 674 difference method, is validated with respect to diffusion and binding tests by means
 675 of comparisons with experimental results. Based on the obtained results, the following
 676 conclusions can be drawn:

- 677 - the proposed model provides a satisfactorily accurate prediction of the diffusivity
 678 of ordinary Portland cement (OPC) with $0.35 \leq w/c \leq 0.60$, especially considering
 679 that no fine tuning of the parameters is performed. In particular, the magnitude
 680 of the predicted diffusivities and their trend vs. the w/c ratio are consistent with
 681 different datasets from the literature;
- 682 - the effect of binding in OPC with $w/c \leq 0.60$ is non negligible and may reduce the
 683 homogenized diffusivity of the HCP by up to 60 %;

- 684 - the heterogeneous nature of the microstructure due to the distribution of the phases
685 in HCP plays a significant role in the diffusion of chlorides, especially including the
686 binding effect. The key role of the percolation properties of the HCP capillary pore
687 system is confirmed;
- 688 - the adoption of the Fickian theory leads to an overestimation of the homogenized
689 diffusivity for very porous HCP (i.e. for $w/c > 0.60$), suggesting a drastic change
690 in the diffusive process in relation to the pore topology;
- 691 - although further validation adopting different cements and w/c ratios is advisable,
692 the first results of simulated and real HCP microstructures show a good agreement
693 provided that for the latter a sufficiently precise estimate of the volume fractions
694 of the phases exhibiting binding is known.

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