



Rejection Mechanism of Ionic Solute Removal by Nanofiltration Membranes: An Overview

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- (1) Input the numerical parameters: number of discretization points, N , grid expansion factor, maximum number of iterations, normalized residuals target and the under-relaxation factor. If the grid expansion factor is higher than one, the distance between consecutive grid nodes expands by this factor from the membrane/solution interfaces to the center of the membrane active layer. If the mesh expansion factor is equal to one, the grid is uniform.
- (2) Input the membrane pore radius, r_p , equivalent membrane thickness, Δx_e , membrane volume charge density, cx and pore dielectric constant, ϵ_p .
- (3) Input the operating parameters: permeate flux, J_v , temperature, T , number of ionic species, N_c , ions diffusivities, $D_{i,\infty}$, concentration of the ions in the feed-solution, $c_{i,b}$, parameters of the mass-transfer correlation, feed Reynolds number, and characteristic length of feed-channel, L .
- (4) Initialize the variables $c_{i,m}$, $c_{i,j}$, $c_{i,p}$, Ψ_j , Ψ_p and ϵ . The concentrations, $c_{i,m}$, $c_{i,j}$ and $c_{i,p}$, are initialized with the bulk concentration, $c_{i,b}$, and the initial values for Ψ_j , Ψ_p and ϵ are taken as zero.
- (5) Compute the coefficients of the coupled linearized system of equations.
- (6) Solve the previous linear system of equations using LU decomposition with partial pivoting. These calculations are accomplished using the subroutine SGEVS contained in the LAPACK package, which is a collection of Fortran subroutines for solving dense linear algebra problems.
- (7) Under-relax the computed solutions.
- (8) Update the coefficients of the coupled linearized system of equations using the under-relaxed variables.
- (9) Compute the normalized residuals.
- (10) Repeat the steps 5–9 until the normalized residuals are below a given target.