

Concentration dynamics in a population model structured in age and a phenotypical trait

Cécile Taing

Laboratoire de Mathématiques et Applications
Université de Poitiers

CEMRACS 2022 : Transport in Physics, Biology and Urban traffic

Context

- Asexual population
- Age structure : birth and death rates depend on the age variable
- Trait structure : competition, adaptation and mutations

The age- and trait-structured population model

$$\left\{ \begin{array}{l} \varepsilon \partial_t n_\varepsilon(t, a, x) + \partial_a n_\varepsilon(t, a, x) + (\rho_\varepsilon(t) + d(a, x)) n_\varepsilon(t, a, x) = 0 \\ n_\varepsilon(t, a = 0, x) = \iint_{\mathbb{R}_+ \times \mathbb{R}^d} \frac{1}{\varepsilon^d} M\left(\frac{x - x'}{\varepsilon}\right) b(a', x') n(t, a', x') da' dx' \\ \rho_\varepsilon(t) = \iint_{\mathbb{R}_+ \times \mathbb{R}^d} n_\varepsilon(t, a, x) da dx, \quad n_\varepsilon(t = 0, a, x) = n_\varepsilon^0(a, x) > 0 \end{array} \right.$$

Goal : To prove $n_\varepsilon(t, a, x) \xrightarrow{\varepsilon \rightarrow 0} \rho(t) p(a, x) \delta_{x=\bar{x}(t)}$

Outline

Age structure

Adding the trait structure

Age- and trait-structured population model

Conclusion

Outline

Age structure

Adding the trait structure

Age- and trait-structured population model

Conclusion

Motivations

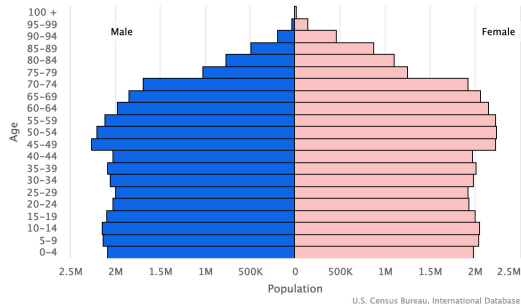


Figure 1: Population pyramid in France in 2020

- demographic : evolution of the age distribution of a human population
- ecological : understanding of the age-dependent birth and death processes of a species

Question : Does the population reach a stable age distribution ?

Age-structured population model

- $a \in \mathbb{R}_+$ age variable
- $n(t, a)$ population density at time t and age a
- $b(a)$, $d(a)$ age-dependent birth and death rates
- $B(t) = \int_{\mathbb{R}_+} b(a)n(t, a) da$ **fertility or renewal term**

Lotka-McKendrick equation

$$\begin{cases} \partial_t n(t, a) + \partial_a n(t, a) + d(a)n(t, a) = 0 \\ n(t, a = 0) = B(t) = \int_{\mathbb{R}_+} b(a)n(t, a) da \\ n(t = 0, a) = n^0(a) > 0. \end{cases}$$

No time-dependency, finite or infinite maximum age, no population migration.

A.G. McKendrick, *Application of mathematics to medical problems* (1926). W. Feller, *On the integral equation of renewal theory* (1941).

Around the age model

On the linear age model

- O. Diekmann, M. Iannelli, B. Perthame, G.F. Webb, ...
- On the stable age distribution : S. Busenberg, M. Iannelli (1985)
P. Michel, S. Mischler, B. Perthame (2005), ...
- On the numerical analysis : J. Douglas Jr., F.A. Milner, D. Sulsky,
A.M de Roos, ...

Extensions of the linear model

- Nonlinear models : logistic term, Allee effect, ...
- Cell division, maturation structure, size structure, ...
- Epidemics : Kermack-McKendrick SIR model
- etc.

Existence and uniqueness

$$S(a) := e^{-\int_0^a d(\sigma) d\sigma} \text{ survival probability at age } a$$

Method of characteristics :

$$n(t, a) = \begin{cases} B(t - a)S(a), & \text{if } a \leq t, \\ n^0(a - t) \frac{S(a)}{S(a - t)}, & \text{if } a > t. \end{cases}$$

Renewal integral equation :

$$B(t) = F(t) + \int_0^t K(a)B(t - a) da,$$

$$\text{with } K(a) = b(a)S(a).$$

Existence and uniqueness of B

There exists a unique continuous solution to the renewal equation

Proof : Picard iterations method

Results : asymptotic behavior

$$n(t, a) \underset{t \rightarrow +\infty}{\sim} e^{\lambda^* t} P(a)$$

with $P(a)$ the stable age distribution

Lotka characteristic equation

$$\widehat{K}(\lambda^*) := \int_0^{+\infty} b(a)S(a)e^{-\lambda^* a} da = 1$$

$\widehat{K}(0)$ is the net reproduction rate

Two approaches :

- Laplace transform of B : λ^* is a simple pole of \widehat{B} .
- Entropy method : λ^* is the principal eigenvalue

$$\int_0^{+\infty} |n(t, a)e^{-\lambda^* t} - \gamma_0 P(a)| \phi(a) da \rightarrow 0$$

Asymptotic behavior : entropy method I

Spectral problem

There exist unique $(\lambda^*, P(a))$ principal eigenlements to

$$\begin{cases} \partial_a P(a) + d(a)P(a) + \lambda^* P(a) = 0, \\ P(0) = \int b(a')P(a') da' = 1, \quad P(a) > 0. \end{cases}$$

From the differential equation,

$$P(a) = P(0)e^{-\lambda^* a - \int_0^a d(\sigma) d\sigma}.$$

Compute λ^* from the renewal term $P(0) = \int bP$,

$$1 = \int_0^\infty b(a)e^{-\lambda^* a - \int_0^a d(\sigma) d\sigma} da.$$

The normalization gives the value of $P(0)$.

Asymptotic behavior : entropy method II

Age profile : $p(t, a) := n(t, a)e^{-\lambda^* t}$

$$\partial_t p(t, a) + \partial_a p(t, a) + d(a)p(t, a) + \lambda^* p(t, a) = 0$$

Then,

$$\partial_t \frac{p(t, a)}{P(a)} + \partial_a \frac{p(t, a)}{P(a)} = 0$$

For $H : \mathbb{R} \rightarrow \mathbb{R}$ convex,

$$\frac{d}{dt} \int \phi(a) P(a) H\left(\frac{p(t, a)}{P(a)}\right) da \leq 0$$

where ϕ is the associated dual eigenvector

$$\begin{cases} -\partial_a \phi(a) + d(a)\phi(a) + \lambda\phi(a) = \phi(0)b(a), \\ \int P(a)\phi(a) da = 1, \quad \phi(a) \geq 0. \end{cases}$$

P. Michel, S. Mischler, B. Perthame, *General relative entropy inequality: an illustration on growth models*, (2005)

Asymptotic behavior : entropy method III

$$\begin{aligned} \partial_t \phi(a) P(a) H\left(\frac{p(t, a)}{P(a)}\right) + \partial_a \phi(a) P(a) H\left(\frac{p(t, a)}{P(a)}\right) = \\ - \phi(0) b(a) P(a) H\left(\frac{p(t, a)}{P(a)}\right) \end{aligned}$$

By Jensen inequality,

$$\frac{d}{dt} \int \phi(a) P(a) H\left(\frac{p(t, a)}{P(a)}\right) da \leq 0$$

Convergence to the stage age profile

Assume $|n^0(a)| \leq c_0 P(a)$. Then,

$$\int_0^{+\infty} |n(t, a) e^{-\lambda^* t} - \gamma_0 P(a)| \phi(a) da \rightarrow 0$$

where $\gamma_0 = \int n^0 \phi$

Outline

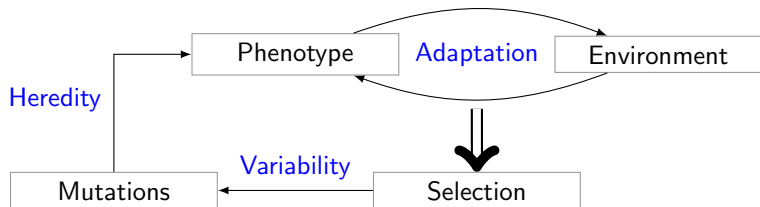
Age structure

Adding the trait structure

Age- and trait-structured population model

Conclusion

Context



- Population structured by a continuous **phenotypic trait** variable
- Interactions with the environment \Rightarrow **non local models**
- **Dirac mass concentration** \iff **selection of the fittest traits**

WKB approach

Time rescaling : $t \leftarrow \frac{t}{\varepsilon}$

$$\varepsilon \partial_t n_\varepsilon(t, x) = n_\varepsilon(t, x) R(x, \rho_\varepsilon(t)) + \varepsilon^2 \Delta n_\varepsilon(t, x)$$

- Hopf-Cole transform:

$$n_\varepsilon(t, x) = e^{\frac{u_\varepsilon(t, x)}{\varepsilon}}$$

- Equation for u_ε :

$$\partial_t u_\varepsilon(t, x) = |\nabla u_\varepsilon|^2 + R(x, \bar{\rho}_\varepsilon(t)) + \varepsilon \Delta u_\varepsilon.$$

- When $\varepsilon \rightarrow 0$, u_ε converges to u **viscosity solution** to the constrained Hamilton-Jacobi equation

$$\begin{cases} \partial_t u(t, x) = |\nabla u|^2 + R(x, \bar{\rho}(t)), \\ \max_{x \in \mathbb{R}^d} u(t, x) = 0, \quad \forall t \geq 0. \end{cases}$$

Biological motivations

- Extensions to various biological problems, e.g. space, migration between patches, nutrients-consumers, by V. Calvez, P.-E. Jabin, S. Mirrahimi, F. Patout, B. Perthame, A. Lam, etc.

Age and trait structure :

- Demographic evolution
- Antagonist pleiotropic theory : some features with positive effects at early ages and negative effects at older ages, and vice versa

Analysis in two cases : without and with mutations

Work done in collaboration with Samuel Nordmann (U. of Tel Aviv) and Benoît Perthame (Sorbonne Université)

Outline

Age structure

Adding the trait structure

Age- and trait-structured population model

Conclusion

Construction of the model

From the renewal equation :

$$\begin{cases} \partial_t n(t, a, x) + \partial_a n(t, a, x) + (d(a, x) + \rho(t)) n(t, a, x) = 0 \\ n(t, a = 0, x) = \iint_{\mathbb{R}_+ \times \mathbb{R}^d} M(x - x') b(a', x') n(t, a', x') da' dx' \end{cases}$$

- $\rho(t) = \int_{\mathbb{R}_+} \int_{\mathbb{R}} n(t, a, x) dx da$ competition
- $M(x - x')$ mutation kernel

Rescaling : $t \leftarrow \frac{t}{\varepsilon}$, $n_\varepsilon(t, a, x) := n(\frac{t}{\varepsilon}, a, x)$

$$\begin{cases} \varepsilon \partial_t n_\varepsilon(t, a, x) + \partial_a n_\varepsilon(t, a, x) + (\rho_\varepsilon(t) + d(x, y)) n_\varepsilon(t, a, x) = 0 \\ n_\varepsilon(t, a = 0, x) = \frac{1}{\varepsilon} \iint_{\mathbb{R}_+ \times \mathbb{R}^d} M\left(\frac{x - x'}{\varepsilon}\right) b(a', x') n_\varepsilon(t, a', x') da' dx' \end{cases}$$

Model without mutations : $M(z) = \delta_0(z)$

$$\begin{cases} \varepsilon \partial_t n_\varepsilon(t, a, x) + \partial_a n_\varepsilon(t, a, x) + (\rho_\varepsilon(t) + d(a, x)) n_\varepsilon(t, a, x) = 0 \\ n_\varepsilon(t, a = 0, x) = \int_{\mathbb{R}_+} b(a', x) n_\varepsilon(t, a', x) da' \end{cases}$$

Factorization and Hopf-Cole transform

$$n_\varepsilon(t, a, x) = p_\varepsilon(t, a, x) e^{\frac{u_\varepsilon(t, x)}{\varepsilon}}$$

Age profile p_ε

$$\begin{cases} \varepsilon \partial_t p_\varepsilon + \partial_a p_\varepsilon + d(a, x) p_\varepsilon + (\partial_t u_\varepsilon(t, x) + \rho_\varepsilon(t)) p_\varepsilon = 0, \\ p_\varepsilon(t, 0, x) = \int_{\mathbb{R}_+} b(a', x) p_\varepsilon(t, a', x) da'. \end{cases}$$

Idea : Constrain $\partial_t u_\varepsilon(t, x) + \rho_\varepsilon(t)$

Factorization

Principal eigenelements

Fix $x \in \mathbb{R}$. There exist unique principal eigenelements $(\Lambda(x), P(a, x))$ to

$$\begin{cases} \partial_a P(a, x) + d(a, x)P(a, x) + \Lambda(x)P(a, x) = 0, \\ P(0, x) = \int b(a', x)P(a', x) da' = 1, \quad P(a, x) > 0. \end{cases}$$

- We define

$$\partial_t u_\varepsilon(t, x) = \Lambda(x) - \rho_\varepsilon(t)$$

- Equation on the age profile $p_\varepsilon = n_\varepsilon e^{-\frac{u_\varepsilon}{\varepsilon}}$

$$\begin{cases} \varepsilon \partial_t p_\varepsilon + \partial_a p_\varepsilon + d(a, x)p_\varepsilon + \Lambda(x)p_\varepsilon = 0, \\ p_\varepsilon(t, 0, x) = \int b(a', x)p_\varepsilon(t, a', x) da'. \end{cases}$$

Convergence results

Convergence of u_ε and p_ε

- $\rho_\varepsilon(t) \xrightarrow[\varepsilon \rightarrow 0]{*} \rho(t)$ in $L^\infty(0, \infty)$
- p_ε converges to a multiple of P for a weighted L^p norm, $p \geq 1$
- u_ε converges locally uniformly to a continuous function u solving

$$\begin{cases} \partial_t u(t, y) = \Lambda(y) - \rho(t), & t > 0, y \in \mathbb{R}^n, \\ \sup_{y \in \mathbb{R}} u(t, y) = 0, & \forall t > 0. \end{cases}$$

With u^0 and Λ concave, and defining $\bar{x}(t) = \operatorname{argmax} u(t, \cdot)$

$$n_\varepsilon(t, a, x) \xrightarrow[\varepsilon \rightarrow 0]{} \rho(t) \frac{1}{\int_x P(a, x)} P(a, x) \delta_{\bar{x}(t)}.$$

Convergence of p_ε

Theorem (convergence of the profile) There exists γ_0 s.t.

$$\int_{\mathbb{R}_+} \left(\frac{p_\varepsilon(t, a, x)}{P(a, x)} - \gamma_0 \right)^2 \mu(a, x) da \xrightarrow{\varepsilon \rightarrow 0} 0, \text{ locally uniformly in } (t, x),$$

where $\mu(a, x) = P(a, x)\Phi(a, x)$ is a probability measure.

Proof. Set $v_\varepsilon(t, a, x) = \frac{p_\varepsilon(t, a, x)}{P(a, x)} - \gamma_0$.

► Compute:
$$\begin{cases} \varepsilon \partial_t v_\varepsilon + \partial_a v_\varepsilon = 0 \\ v_\varepsilon(t, a = 0, x) = \int_a v_\varepsilon b P \end{cases}$$

► Introduce the **dual eigenfunction** $\Phi(a, x)$ of $P(a, x)$.

► Entropy
$$E_\varepsilon(t, x) = \int_{\mathbb{R}_+} v_\varepsilon^2 P(a, x) \Phi(a, x) da.$$

►
$$\partial_t E_\varepsilon(t, x) \leq 0 \implies 0 \leq E_\varepsilon(t, x) \leq E_\varepsilon(t = 0, x) \implies E_\varepsilon(t, x) \xrightarrow{\varepsilon} 0.$$

Convergence of u_ε

Constraint on u

u_ε converges locally uniformly towards u , solution to

$$\begin{cases} \partial_t u(t, x) = \Lambda(x) - \rho(t) \\ \sup_{x \in \mathbb{R}} u(t, x) = 0, \forall t \geq 0, \end{cases}$$

- $0 < \underline{\rho} \leq \rho_\varepsilon(t) \leq \bar{\rho}$, $\rho_\varepsilon(t) \xrightarrow[\varepsilon \rightarrow 0]{*} \rho(t)$, in L^∞ (after extraction)

$$\text{Convergence: } u_\varepsilon(t, x) \xrightarrow[\varepsilon \rightarrow 0]{} u(t, x) = u^0(x) + t\Lambda(x) - \int_0^t \rho.$$

- For all $t \geq 0$, $\sup_{x \in \mathbb{R}} u(t, x) = 0$,

$$\int_0^t \rho = \sup_y [u^0(x) + t\Lambda(x)]$$

Numerical simulations

$$A(a, x) = 1, \quad b(a, x) = \frac{10x}{1 + a^2}, \quad d(a, x) = x^3(2 + a/3),$$

$$p^0(a, x) = \exp(-0.8a), \quad u^0(x) = -\frac{(x - 0.5)^2}{2}.$$

Implicit-explicit finite differences scheme

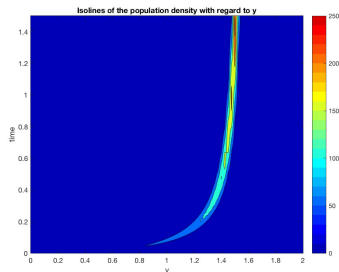
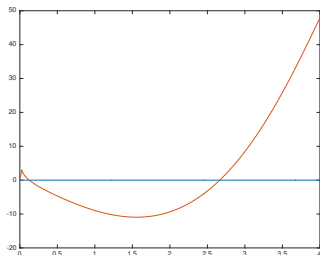


Figure 2: (Left) effective fitness $\Lambda(x)$, (right) total population $t \mapsto \rho_\varepsilon(t)$

Numerical simulations

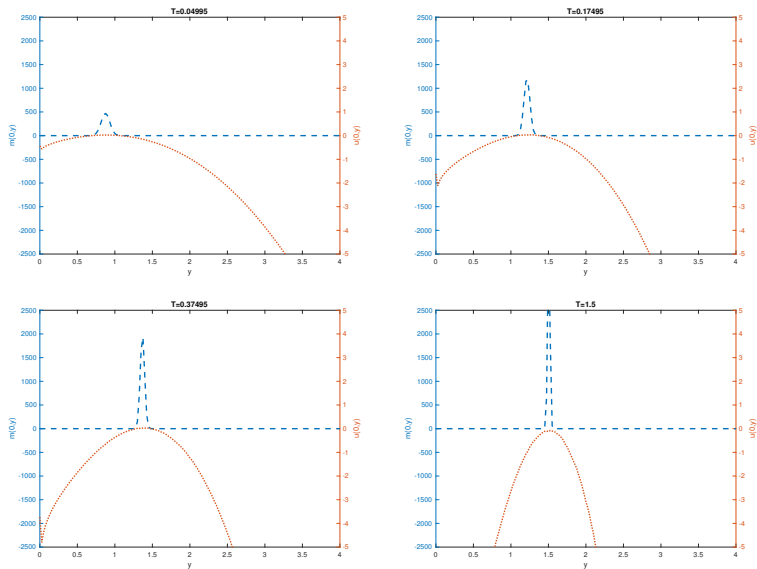


Figure 3: (blue) population of age 0, (red) $u(t, x)$

With mutations : formal computations

- Same approach: $n_\varepsilon(t, a, x) = p_\varepsilon(t, a, x) e^{\frac{u_\varepsilon(t, x)}{\varepsilon}}$,

$$\begin{cases} \varepsilon \partial_t p_\varepsilon + \partial_a p_\varepsilon + d(a, x) p_\varepsilon = (-\partial_t u_\varepsilon(t, x) - \rho_\varepsilon(t)) p_\varepsilon, \\ p_\varepsilon(t, 0, x) = \iint_{x', a'} \frac{1}{\varepsilon^d} M\left(\frac{x-x'}{\varepsilon}\right) e^{\frac{u_\varepsilon(t, x') - u_\varepsilon(t, x)}{\varepsilon}} b(a', x') p_\varepsilon(t, a', x') \end{cases}$$

- Change of variable $x' \mapsto z = \frac{x' - x}{\varepsilon}$

$$\begin{aligned} p_\varepsilon(t, 0, x) &= \int_z M(z) e^{\frac{u_\varepsilon(t, x + \varepsilon z) - u_\varepsilon(t, x)}{\varepsilon}} \int_{a'} b(a', x + \varepsilon z) p_\varepsilon(t, a', x + \varepsilon z) da dz \\ &\rightarrow \left(\int_z M(z) e^{\partial_x u(t, x) z} dz \right) \left(\int_{a'} b(a', x) p(t, a', x) da \right) \end{aligned}$$

- Formal limit

$$\begin{cases} \partial_a p + d(a, x) p = (-\partial_t u(t, x) - \rho(t)) p, \\ p(t, 0, x) = \int_z M(z) e^{\partial_x u(t, x) z} dz \int_{a'} b(a', x) p(t, a', x) da \end{cases}$$

Hamiltonian

From $\partial_a p + d(a, x)p = (-\partial_t u_\varepsilon(t, x) - \rho_\varepsilon(t))p$,

$$p(t, a, x) = p(t, 0, x)e^{-\int_0^a d(\alpha)d\alpha - (\partial_t u(t, x) + \rho(t))a}.$$

From the renewal term,

$$\frac{1}{\int_{\mathbb{R}} M(z)e^{\partial_x u(t, x)z} dz} = \int_0^{+\infty} b(a, x)e^{-\int_0^a d(\alpha)d\alpha - (\partial_t u(t, x) + \rho(t))a} da,$$

$$\widehat{K}(\partial_t u(t, x) + \rho(t)) = \left(\int_z M(z)e^{\partial_x u(t, x)z} dz \right)^{-1}$$

Hamilton-Jacobi equation :

$$\partial_t u(t, x) = H(\partial_x u(t, x)) - \rho(t), \quad \widehat{K}(H(p)) = \left(\int_z M(z)e^{pz} dz \right)^{-1}$$

V. Calvez, P. Gabriel, A. Gonzalez Mateos, *Limiting Hamilton-Jacobi equation for the large scale asymptotics of a subdiffusion jump-renewal equation*, (2019)

Convergence results

Spectral problem

Find $\Lambda(x, \eta)$, $P(a, x, \eta)$ solution to

$$\begin{cases} \partial_a P(a, x, \eta) + d(a, x)P(a, x, \eta) + \Lambda(x, \eta)P(a, x, \eta) = 0, \\ P(0, x, \eta) = \eta \int b(a', x)P(a', x, \eta) da' = 1, \quad P(a, x, \eta) > 0. \end{cases}$$

Results :

- Construction of an approximated problem

$$\begin{cases} \varepsilon \partial_t p_\varepsilon + \partial_a p_\varepsilon + d(a, x)p_\varepsilon = (-\partial_t u_\varepsilon(t, x) - \rho_\varepsilon(t)) p_\varepsilon, \\ p_\varepsilon(t, 0, x) = \eta_\varepsilon(t, x) \int_{a'} b(a', x)p_\varepsilon(t, a', x) dx \end{cases}$$

- Convergence of $U_\varepsilon(t, x) = u_\varepsilon(t, x) + \int_0^t \rho_\varepsilon$

Outline

Age structure

Adding the trait structure

Age- and trait-structured population model

Conclusion

In conclusion

Results :

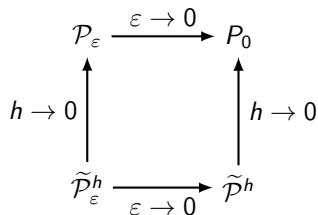
- Factorization : $n_\varepsilon(t, a, x) = p_\varepsilon(t, a, x)e^{\frac{u_\varepsilon(t, x)}{\varepsilon}}$
 - ▶ trait profile $e^{\frac{u_\varepsilon(t, x)}{\varepsilon}}$ led by $\Lambda(x)$
 - ▶ age profile $p_\varepsilon \rightarrow \gamma_0 P$
- Without mutations : convergence with entropy estimates
- With mutations : Hamilton-Jacobi equation on $u + \int_0^t \rho$
 - ▶ Definition of an approximated problem \tilde{P}_ε
 - ▶ Construction of \tilde{u}_ε and \tilde{p}_ε
 - ▶ Hamilton-Jacobi equation on $\tilde{u} + \int_0^t \rho$

Recent results

Convergence of p_ε : Nordmann, Perthame (2021)

- estimates of U_ε in $W_{loc}^{1,r}$ with $1 \leq r < \infty$
- $p_\varepsilon \xrightarrow[\varepsilon \rightarrow 0]{L^\infty^*} \gamma_0 P$

Asymptotic preserving scheme: ongoing work with H. Hivert



Based on the spectral problem : **not working**

- compute Λ and P
- compute U_ε and \tilde{p}_ε with extrapolation on η_ε
- $p_\varepsilon = \tilde{p}_\varepsilon + O(\varepsilon)$

Need of another approach

Thank you for your attention !